

STIC Search Report Biotech-Chem Library

STIC Database Tracking Number:

TO: Karen A Lacourciere

Location: rem/2d15/2c18

Art Unit: 1635

Monday, March 08, 2004

Case Serial Number: 09/848868

From: Alex Waclawiw

Location: Biotech-Chem Library

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Phone: 308-4491

Alexandra.waclawiw@uspto.gov

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CC The present invention relates to anti CC mRNA sequence for a selected protein, CC 1 to about 25 base pairs downstream o CC in the preprocessed mRNA. The antisen	aim 1	lice-region antisense	XX DR WPI; 2002-066533/09.	PI Iversen PL, Hudziak R;	PA (AVIB-) AVI BIOPHARMA INC.	PR 04-MAY-2000; 2000US-0202376P	PF 04-MAY-2001; 2001WO-US014410	PD 08-NOV-2001.	PN WO200183740-A2.	OS Homo sapiens.	nsense; splice region	man p53 coding sequenc	DT 15-MAR-2002 (first entry)	AC AAS19106;	ID AAS19106 standard; DNA; 18 BP.	RESULT 1 AAS19106		200 14 77.8 113604 8	14 77.8 67191 9 14 77.8 113604 8	196 14 77.8 61457 9	194 14 77.8 8131 9	92 14 77.8 5389 4 93 14 77.8 7065 7	90 14 77.8 3294 4 91 14 77.8 5344 4	14 77.8 3294 4	87 14 77.8 2679 5	14 77.8 2679 5 14 77.8 2679 5	84 14 77.8 1/66 4 87 14 77.8 2453 5	14 77.8 549 4	180 14.4 80.0 349980 3 181 14 77.8 469 8	79 14.4 80.0 349980 3	177 14.4 80.0 110000 3 178 14.4 80.0 129722 6	175 14.4 80.0 73465 6 176 14.4 80.0 110000 3	73 14.4 80.0 56485 3 74 14.4 80.0 65632 3	.4 80.0 24329 4 .4 80.0 24329 4	14.4 80.0 24279 4
relates to antisense compositions targeted to an lected protein, at a region having its 5' end from rs downstream of a normal splice acceptor junction NA. The antisense compound is Rnase-inactive, and	English.	composition and method.				••	3.				n; mRNA splice processing inhibition; expression inhibition; human; HIV-1;	equence SEQ					ALIGNMENTS		ADB72641 Ac ADA42607 Ac												'4 4'	01		75856 Aak75856 75000 Aak75000	

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Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This invention describes a vector containing a recombinase gene which i controlled by (1) a promoter and action of which is in turn dependent a specific transcription factor and (2) a desired gene to be expressed and two target sequences of the recombinase. Also described in the invention are (i) a host cell for introduction of the vector and (ii) a in vitro technique in which killing of cells without the specific transcription factor is selectively performed after the vector is introduced into a host cell by an in vitro technique. The technique is used for gene therapy e.g. in cancer treatment. The products of the invention allow the use of a specific vector specifically and safely, there is little toxicity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gene expression specific to cells free from specific transcription factor with constructed recombinase expression unit after infecting cells - allowing expression of target gene in gene therapy, particularly in cancer treatment.
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                                      Homo
                                                                                            L-oncogene;
                                                                                                                                   Human breast
                                                                                                                                                                       14-AUG-1998
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                                                                                                                                                                                                                                                 AAV17739 standard; cDNA; 140
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 37 BP; 7 A; 13 C; 9 G; 8 T; 0 U; 0 Other;
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                                                                                            diagnosis; treatment; ovarian cancer; antibody; antisense;
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                                                                                                                                 cancer related gene BC532L.
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74245
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WPI; 199
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 2; Fig 1c; 96pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human breast cancer related genes - used for diagnosis, detection and therapy of breast and ovarian cancers.
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                                                                                                                                                                BC534; BC538;
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02-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                   Sequence 140 BP; 29
                          Bergmann JE,
                                            (BERG/)
                                                                                         04-MAR-1994;
                                                                                                           29-SEP-1994.
                                                                                                                            WO9421791-A1
                                                                                                                                                                                 p53 antisense
                                                                                                                                                                                                                              AAQ74245;
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                                                                       16-MAR-1993;
         1994-357732/44.
                                                                                                                                             sapiens
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PREDDIE E R.
 AAR51627.
                                                     BERGMANN J E.
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(first entry)
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                           Preddie RE;
                                                                                                                                                                                 strand cDNA encoding protein BC538.1.
                                                                                                                                                               BC538.1; p53 antisense strand; breast cancer;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New p53 anti sense proteins - used to develop prods, for the diagnosis, prediction and treatment of breast cancer and related cancers.
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                                                                                                                          WPI; 1998-169156/15.
P-PSDB; AAW48347.
                                                                                                                                                                                                                                 22-AUG-1997;
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                                                                                                                                                          Bergmann JE,
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Human breast cancer related genes, 1-oncogenes (AAV17738-V17753), a proteins encoded can be used in the diagnosis and treatment of brea ovarian cancer. Antibodies and multiple antigenic peptide epitopes used to detect the presence of the proteins. The antibodies can als used to block the activity of the proteins. Antisense molecules can used to prevent expression of the proteins

17753), and the t of breast and epitopes can be

pes can k also be

Claim 1; Fig 1e; 96pp; English. Human breast cancer related genes - detection and therapy of breast and

used for diagnosis, pre-symptomatic ovarian cancers.

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T59673/
                                   A novel method has been produced for analysing a target nucleotide sequence. The target sequence having a first segment, a second segment and a third segment between the first and second, which is formed of at least one, but less than four different, nucleotide(s) where the third segment has a nucleotide or nucleotide sequence in a first state or different second state. The present sequence is a partial human p53 gene sequence which is used as an example of a target nucleotide sequence. The method can be used to analyse the entire sequence (portion) of a known gene and to analyse infectious disease or to determine if a sample is from a particular source, such as for criminal investigations or parental determination. Especially the method is used to determine the existence or absence of mutations consisting of one or a few nucleotides
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                                                                                                                                                                                                                                                                                                                                     Disclosure; Page 45-46; 78pp; English.
                                                                                                                                                                                                                                                                                                                                                                                Analysis of target nucleotide sequence which exits in a first state different second state - useful for determination of point mutation
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       07-JUN-1995;
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/note=
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Pred. No. 25;
0; Mismatches
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186 T;

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Query Match Best Local Similarity "Atches 18; Conserve

Conservative

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Mismatches

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Indels

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Gaps

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100.0%;

Score 18; Pred. No.

7;

Length 1024;

Sequence 1024

BP;

225 A; 317

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263 G; 219 T; 0 U; 0 Other;

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RESULT 8
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ID ABZ88
XX ABZ88767//
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ABZ887
                                     CC The present invention occurred.

CC The present invention

CC toxicological response to an agent, which comprises comparing the

CC expression profile of one or more human toxic response genes to a

CC reference gene expression profile indicative of toxicity, and so

CC determining the presence of a toxic response to the agent. Also

CC described: (1) an array comprising one or more polynucleotides selected

CC from the genes corresponding to the partial sequences given in ABS82842

CC to ABS84744, or their fragments of at least 20 nucleotides, or homologues

cc response gene plays a role on toxic response pathways by determining the

CC expression profile of the gene after exposure of cells or a human subject

to a known toxic pharmaceutical or industrial agent, comprising: (a)

CC was exposed to an agent (b) obtaining the test gene expression profile

CC was exposed to an agent; (b) obtaining the test gene expression profile

CC profile to the expression profile of a gene with a similar function or

CC comparing the test profile to the expression profile of that gene after

exposure to other known toxic compounds. The methods are useful for

cc profile to the response to a cellular, organ

cc rystem level. The arrays comprising the human genes are useful for

cr system level. The arrays comprising the human genes are useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Determining a toxicological response to an agent, useful for screening drugs, comprises comparing the expression profile of one or more human toxic response genes to a reference gene expression profile indicative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Alen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 1; Page
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Toxicologically relevant human nucleotide sequence #926
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABZ83767
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention describes a method toxicological response to an agent, which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             toxicity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2003-268322/26
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PHASE-1 MOLECULAR TOXICOLOGY INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
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27;
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RESULT 9
AAZ52304/c
ID AAZ52304 standard; c
XX AAZ52304;
XX AAZ52304;
XX AAZ52304;
XX AAZ52304;
XX AAZ52304;
XX AAZ52304;
XX C4-JUL-2000 (first
XX Human p35 (p53 isoform; p
XX Human p53 isoform; p
XX C20-APR-2000.
PH Key C200022127-A1.
XX C20-APR-2000.
XX C20-APR-200
RESULT 10
AAX04533/c
ID AAX045
XX
AC AAX045
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                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                          Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present cDNA sequence encodes the human p53 isoform, p35. This p53 isoform is truncated to eliminate a substantial portion of the C-terminal end of the protein. The deleted portion correspond to those encoded by exon 7 and by exons corresponding to amino acids lying C-terminal to exon 7, p35 is useful as a marker of myocardial infarction (by the indication of hypoxia) and in the control of cell proliferation both in vivo and in vitro. It was able to increase the transactivation achieved through cotransfection with wild type p53. The combination of p35 with wild type p53 produce an enhanced inhibition of primary incorporation at 24 hours
  AAX04533
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 18; Fig 4; 25pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel isoform of p53 useful as a marker controlling cellular proliferation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2000-317984/27.
P-PSDB; AAY70714.
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                                                   AAX04533 standard; DNA; 1303
                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 1070 BP; 233 A; 338 C; 282 G; 217 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            compared to wild type p53 alone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (BGHM ) BRIGHAM & WOMENS HOSPITAL
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                                                                                                                                                                                                                                                                                                Conservative
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/product= "Human p35 protein"
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                                                                                                                                                                                                                                                                                                                       Score 18;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ВР
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RESULT 11
ABS737/c
ID ABS733
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OHO DE DNA en
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Chromo
KW prolif
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present sequence encodes a human p53 protein. The protein is used in the assay of the invention. The specification describes an assay method for a compound able to modulate the interaction between ATM or a protein having an associated kinase activity and p53 or a protein having homologous phosphorylation sites. The assay comprises contacting a peptide fragment ATM with a relevant fragment of p53 and a test compound, and determining the interaction or binding between the substances and the test compound. The assay method is useful for screening for compounds able to modulate the interaction between ATM and p53. The screened agents, peptide fragments and nucleic acids are useful for therapy involving modulating ATM action e.g. in the treatment of cancer, immunosuppression or HIV infections by modulating phosphorylation of p53 by ATM, and for purifying the proteins ATM and ATR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             p53;
Chromosome aberration; oncogenic fusion protein; cancer; oncogene; proliferative disease; cellular protein isoform; heat shock protein 90;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Assay method for compounds modulating the interaction of ATM and useful for the treatment of e.g. cancer, immunosuppression and H infections and for the purification of the proteins ATM and ATR.
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                                                                                                      04-DEC-2002
                                                                                                                                                  ABS73327;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 1303 BP;
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P-PSDB; AAW84270.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ataxia telangiectasia; ATM protein; assay; interaction; kinase activity;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DNA encoding human p53 protein
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                                                                                                                                                                                         standard; DNA; 1303
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                                                                                                                                                                                                                                                                                                  CCCGGAAGGCAGTCTGGC
                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                           human oncogene p54
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         292 A; 403 C; 348
                                                                                                      entry)
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Pred. No.
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RESULT 12
ACC49400/c
ID ACC494
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AC ACC494
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DT 24-JUN
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Human
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KW Human;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CC The invention describes a method of treating genetically-defined disease CC associated with chromosomal aberrations yielding oncogenic fusion CC proteins (1), treating cancerous cells containing (1) in a heterogeneous CC protein (1), treating cancerous cells containing (1) in a heterogeneous CC cell population, treating proliferative disease associated with mutant CC protein or cellular protein isoforms (II) dependent on heat shock protein CC described disease with chromosomal aberration yielding CC genetically-defined disease with chromosomal aberration yielding CC quotein in heterogeneous cell population, treating proliferative disease (e.g. rheumatoid arthritis or cancer) associated with mutant protein or CC cellular protein isoform dependent on heat shock protein (HSP)-90 (e.g. 53), or selectively treating cells expressing mutant protein or cellular protein isoform dependent on heat shock protein (HSP)-90 (e.g. protein isoform dependent on heat shock protein (HSP)-90 (e.g. protein isoform dependent on heat shock protein described is useful CC or a disease characterised by a solid tumour such as T or B cell CC imphoma, chronic myeloid leukaemia (CML), APL, ALL, AML, NHL and CMML, CC or a disease characterised by a solid tumour such as papillary thyroid CC arctinoma, Ewing's sarroma, melanoma, liposarcoma, rhabdomyosarcoma and CC carcinoma, The method is also useful for treating viral infections. This sequence encodes a human oncogenic protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Treating genetically-defined disease associated with chromosomal aberrations yielding oncogenic fusion proteins, e.g. cell proliferative diseases, involves administering an inhibitor of heat shock protein 90.
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Human; ATM; p53; ATR; FRP1;
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                                                               Human
                                                                                                                                     24-JUN-2003
                                                                                                                                                                                                                                                                    ACC49400 standard;
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                                                                  p53 encoding cDNA.
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FRAP related protein; DNA-PK;
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Pred. No. 28;
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113

CCCGGAAGGCAGTCTGGC

18 96 Query Match Best Local Matches 1

Similarity

100.0%;

Score 18; D Pred. No. 28 ; Mismatches

7;

Length 1303; ; Indels

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Gaps

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The present invention describes an assay (MI) for a compound able to a ffect DNA binding by an ataxia-telangiectasia mutated (ATM) protein or a comprotein having an associated kinase activity. MI comprises bringing into contact a substance which is ATM or a protein having an associated kinase activity which is able to bind DNA and a test compound, and determining comparibed: (1) an agent (I) capable of affecting DNA binding by ATM contained using MI; (2) purifying (M2) ATM or TATR; and (4) a substantially compared the compound activities. MI is useful for assaying for a compound able to affect DNA binding by ATM as anti-HIV, cytostatic, antipsoriatic cardinated using binding by ATM or ATM has anti-HIV; cytostatic, antipsoriatic compound activities. MI is useful for assaying for a compound able to affect DNA binding by ATM or a protein having an associated kinase continuity. (I) is useful for assaying an associated kinase continuity. (I) is useful in therapy involving modulating ATM action or in the manufacture of a medicament for modulating ATM action. (II) is useful for treating by mumans with a taxia celangicetasia, AIDS or cancer, for treating or preventing disease states associated with premature and normal aging for regulating immune cycle check point arrest in the absence of cellular damage, which may be useful check point arrest in the absence of cellular damage, which may be check point arrest in the absence of cellular damage, which may be considered encodes human p53 which is given in the exemplification of the present overtion.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DNA dependent protein kinase catalytic subunit; anti-HIV; cytostatic; antipsoriatic; antitumour; ataxia-telangiectasia; AIDS; cancer; tumour; psoriasis; hyperproliferative disorder; chromosome 17p13.1; gene; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Assay for compound affecting DNA binding by ataxia-telangiectasia mutated gene, by bringing into contact the gene, protein with kinase activity, DNA and test compound, and determining binding of the gene and DNA.
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P-PSDB; ABP97119.
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Sequence 1303 BP; 289 A; 401 C; 344 G; 257 T; 0 U; 12 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Fig 7bi-ii; 129pp; English.
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16-JUL-1998;
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98GB-00015423.
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note= "illegible parts of the nucleotide
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AAD00088;

AAD00088 standard; DNA; 1307

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RESULT 13
AADOOOB6/c
ID AADOOO
XX AADOOO
XX AADOOO
XX AADOOO
XX Human;
XX Human;
XX Human;
XX Homo (
XX Homo (
XX Homo (
XX Homo)
XX Homo
XX 
RESULT 14
ABX15394/c
ID ABX153
XX
AC ABX153
XX
DT 01-MAY
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                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The patent discloses a method for the treatment of systemic lupus erythematosus (SLE) by down-regulating the autoimmune response to the C-terminal DNA-binding domain of p53 protein by an active compound comprising C-terminal DNA-binding domain of p53, monoclonal antibodies (Ab1) specific to this domain, monoclonal antibodies (Ab2) specific to Ab1 and peptides based on the complementarity determining region of heavy and light chain of Ab1 and Ab2. The active compound is useful in the diagnosis, prevention and treatment of SLE in humans. The present sequence is a human tumour-associated antigen p53 DNA. Antibodies against the C-terminal DNA-binding domain of the p53 protein can be raised and used for diagnosis and treatment of SLE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human tumour-associated antigen p53 DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Treatment of systemic lupus erythematosus by down-regulating the autoimmune response to the C-terminal DNA-binding domain of the p53 protein by an active compound comprising of antibodies to p53 or fragments of p53.
     01-MAY-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 1307 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Page 79-81; 87pp; English.
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                                                       ABX15394;
                                                                                                         ABX15394 standard;
                                                                                                                                                                                                                                                                                                                                                                                     Local
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                                                                                                                                                                                                                                                    117
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18; Conserv
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                                                                                                                                                                                                                                                    CCCGGAAGGCAGTCTGGC 100
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                                                                                                                                                                                                                                                                                                                                                             Conservative
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  (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    293 A; 404 C; 350
                                                                                                         DNA; 1307
                                                                                                                                                                                                                                                                                                                                                                               100.0%;
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Pred. No. 28;
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       260 T;
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Human; p53; degradation; cervical cancer; tumour; ovarian cancer; glioma carcinoma; squamous cell carcinoma; lung cancer; pancreatic cancer; leukaemia; lymphoma; neuroblastoma; sarcoma; osteosarcoma; glioblastoma; colon carcinoma; melanoma; choriocarcinoma; breast carcinoma; gene; ds; neuroblastoma; rhabdomyosarcoma; Mdm2; cytostatic; antitumour.
                                                                                                                                                                                                                                                                                                                                                                                        Human p53 DNA.
                                                                                                                                                                                                                                                                                                                                Homo sapiens.
                                                                                                                                                                                                                                 08-DEC-1999;
                                                                                                                                                                                                                                               07-DEC-2000; 2000US-00732384
                                                                                                                                                                                                                                                              19-SEP-2002.
                                                                                                                                                                                                                                                                             US2002132977-A1.
                                                                                                                                                                                                          (YUAN/) YUAN (GUJJ/) GU J.
                                                                                                                                                                                                                  2
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                                                                                                                                                                                                                                                                                           /product= "Human p53"
                                                                                                                                                                                                                                                                                                                                                                           glioma;
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Novel polypeptide that inhibits degradation of endogenous mammalian cell, useful for treating cancer, e.g. cervical tumor such as sarcoma or carcinoma. p53 in cancer or a

P-PSDB; ABG73431.

2003-197937/19.

Yuan Z,

Gu J

Example 2; Page 3; 21pp; English.

CC degradation of endogenous p53 in a mammalian cell. The polypeptide or a cc synthetic polypeptide comprising a region of the pure polypeptide or a cc synthetic polypeptide comprising a region of the pure polypeptide is cuseful for inhibiting degradation of endogenous p53 in a mammalian cell, ce g; in cervical cancer cells or tumour cells selected from sarcoma, cc carcinoma, squamous cell carcinoma, ovarian cancer, lung cancer, cc pancreatic cancer, leukaemia, lymphoma, glioma, neuroblastoma, corteosarcoma, colon carcinoma, melanoma, choriocarcinoma, breast carcinoma, glioblastoma, neuroblastoma and rhabdomyosarcoma cells. The cc carcinoma, glioblastoma, neuroblastoma and rhabdomyosarcoma cells. The cc cell comprises an Mdm2 amplification or overexpresses Mdm2. A degradation cresistant p53 polypeptide is useful for inhibiting tumour growth, where the tumour is a cervical cancer and comprises ap53 mutation. The cc polypeptide, and DNA encoding the polypeptide, are useful for treating cc cancer. This sequence represents DNA encoding the human p53 polypeptide

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Sequence 1307 BP; 293 A; 404 C; 350 G; 260 T; 0 U; 0 Other;
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멂 7;

Length 1307;

0

밁 S Query Match Best Local S Matches 18 117 18; \vdash Similarity CCCGGAAGGCAGTCTGGC 18 CCCGGAAGGCAGTCTGGC 100 Conservative 100.0%; 0, Score 18; Pred. No. Mismatches 0 Indels ٥, Gaps

RESULT 15
AAT32831/c
ID AAT32
XX
AC AAT32
XX
DT 06-NC
XX
DE Humas AAT32831; AAT32831 standard; DNA; 1317 ВP

Human p53 tumour suppressor protein gene

06-NOV-1996

(first entry)

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RESULT 16
AAV18151/c
ID AAV18151 standard; DI
XX
AC AAV18151;
XX
DT 21-AUG-1998 (first 6
XX
DE Nucleotide sequence 6
XX
KW Human p53 gene; vacc;
XW mutation; antigen; cc
XX
CS Homo sapiens.
XX
FH Key Locat
FT CDS 136.
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Best Local S
Matches 18
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28-APR-1995;
01-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                   A DNA sequence (AAT32831) codes for wild-type p53 protein (AAW02617), a sequence-specific DNA binding protein which has tumour suppressor function. p53 regulates cell proliferation and apoptosis and participates in cellular response to DNA damaging agents. It is inactivated in more than half of all human tumours. DNA constructs encoding p53 proteins with altered tetramerisation domains that retain wild-type p53 function can be used in gene therapy to treat abnormal cell proliferation, esp. cancer, or to induce immune tolerance to facilitate transplants and treat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         p53 protein; tumour suppressor; tetramerisation domain; chimaeric protein; gene therapy; vector; cell proliferation; cancer; apoptosis; autoimmune disease; immune tolerance; ds.
                                                                                                                                                                                                                                                                                                                                                                                Sequence 1317 BP;
                                                                                                                                                                                                                                                                                                                                                                                                            autoimmune diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Page 66-68; 123pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New chimaeric p53 protein with heterologous tetramerisation domain - and related DNA and vectors, useful for treating abnormal cell proliferation, esp. cancer, auto-immune disease, etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Halazonetis TD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              27-NOV-1995;
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                                                                                                                                                                                                                                                                           CCCGGAAGGCAGTCTGGC 110
                                                                                                                                                                                                                                                                                                 ccceeaaeccaercreec 18
                                                                                             gene; vaccine; chemokine; cytokine;
                                                                                                                        sequence
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95US-00431357.
95US-00456623.
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136. .1317
/*tag= a
Location/Qualifiers
136. .1317
/*tag= a
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                                                                                 cancer; ds.
                                                                                                                                                                                                      DNA; 1317
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                                                                                                                                                  entry)
                                                                                                                     human p53 gene.
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Pred. No.
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                                                                                                immune
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                                                                                              response; tumour;
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RESULT 17
AAV21414/c
ID AAV214
XX AAV214
XX AAV214
XX Human
XX Probe;
XW in sit
XX Synthe
OS Homo s
YX WO9808
XX Synthe
OS Synthe
OS Synthe
OS Homo s
YX Hom
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  of the invention inclving a pharmaceutical or vaccine comprising DNA encoding p53 and optionally a chemokine or cytokine. The compositions methods can be used for inducing immune responses to tumour cells whice over-express p53 or express mutated p53. They can be used to induce a response to tumour antigens to cause a regression of existing tumours and/or prevent the development of cancers, in e.g. high risk patients
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pharmaceutical or vaccine comprising DNA encoding p53 and optionally chemokine or cytokine - is useful for treatment and prevention of tumours by inducing protective immune response.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 1317 BP; 297 A; 408 C; 350 G; 262 T; 0 U; 0 Other;
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                           WPI; 1998-179448/16
                                                                         Kigawa K,
                                                                                                                                                                             29-AUG-1996;
26-DEC-1996;
                                                                                                                                                                                                                                                        29-AUG-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic.
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P-PSDB; AAW46858.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    gene mapping; RecA-like recombinase; human p53; lambda phage;
u hybridisation; infectious disease; gene therapy; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  l Similarity
18; Conserv
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                                                                         Yamanaka M,
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                                                                                                                                                                             96JP-00229061
96JP-00347090
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Pred. No.
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RESULT 18
AAV446
AC AAV46
AC AAV446
AC AAV46
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hydrophobic; p53 protein; mutant; oligomerisation domain; dimerisation; therapeutic agent; biotechnology; tumour suppressor; apoptosis; human; transplant rejection; autoimmune disease; systemic lupus erythematosus; rheumatoid arthritis; psoriasis; atherosclerosis; arterial restenosis; abnormal cell proliferation; interleukin-2 receptor complex; three-dimensional structure; 3D structure; transmembrane receptor; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  19-OCT-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human wild-type p53 protein encoding DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAV44695
   WPI; 1998-414033/35
P-PSDB; AAW69217.
                                                                                            Halazonetis TD
                                                                                                                                                                                                                                                                                                                                                                                                   WO9831703-A1
                                                                                                                                                       (WIST-) WISTAR
                                                                                                                                                                                                                  17-JAN-1997;
                                                                                                                                                                                                                                                                                15-JAN-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        standard; DNA; 1317
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                                                                                                                                                                                                                  97US-0035458P
                                                                                                                                                                                                                                                                                98WO-US000853
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /product= "wild-type p53 protein'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            293 A; 407 C; 359
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                                                                                                                                                          ANATOMY & BIOLOGY
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Pred. No.
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properties.
                                                                                                                                                                              Altering three-dimensional structure of protein without denaturing it by replacing large hydrophobic amino acids with small ones, or vice versa, used to, e.g. produce p53 proteins with altered oligomerisation
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Example 1; Page 33-34; 49pp; English

CC denaturing the protein The method comprises identifying hydrophobic corresidues in the protein and classifying the residues as large or small corresidues have been substituted are produced and they can be analysed for residues have been substituted are produced and they can be analysed for cresidues having a hydrophobic core. A p53 oligomerisation domain can be caltered by this method. Vectors containing nucleic acids encoding a p53 oligomerisation domain and fusion proteins of the p53 oligomerisation domain and fusion proteins of the p53 oligomerisation can be considered by this method. Vectors containing nucleic acids encoding a p53 oligomerisation domain and fusion proteins of the p53 oligomerisation containing nucleic acids encoding a p53 oligomerisation domain and fusion proteins of the p53 oligomerisation of diagnostic agents, in biotechnology and other industrial applications. Typically altered p53 is useful as tumour suppressor, to induce apoptosis of proliferating lymphocytes, to prevent transplant rejection, to treat curoimmune diseases such as systemic lupus erythematosus and rheumatoid carthritis, and to suppress proliferation in cases of psoriasis, catheroscists and arterial restencial respectual proliferation. Typical of interleukin-2 receptor complex (for screening drugs that bind the native receptor and as therapeutic decoys) and transmembrane receptors. The p53 oligomerisation domain may also be used to induce dimerisation of DNA-control but have better stability, binding and lower molecular weight (so enter cells more easily). When expressed from a gene therapy vector, the altered p53 will not be sequestered into inactive heterodimers by mutative heterodimers by mutative heterodimers by mutative heterodimers by This DNA encodes or altering present in the cells the three-di wild-type p53 protein. The invention provides a method hree-dimensional (3D) structure of a protein, without

Sequence 1317 BP; 297 A; 408 C; 350 G; 262 T; 0 U; 0 Other;

S Matches Query Match Best Local (Local Similarity 18; 1 CCCGGAAGGCAGTCTGGC Conservative 100.0%; 18 0; Score Pred. Mismatches No. ب ب <u>.</u> Length 1317; Indels 0 Gaps 0

RESULT 19 AAC84597/G ID AAC844597/G ID AAC84 XX AAC84 XX AAC84 XX D2-AE XX S-PHA XX S-PHA XX CUL-1 KW Bad; XX Homo XX AAC84597

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127

CCCGGAAGGCAGTCTGGC

standard; DNA; 1317 В₽

02-APR-2001 (first entry

Human p53 protein encoding DNA.

S-phase kinase associated protein; SKP1; SKP2; SKP2-like protein; ZF; CUL-1; cullin; CDC53; p27; cyclin E; Max; Mad; c-Myc; MDM2; p53; Bax; Bcl-2; tumour; cytostatic;

Homo sapiens.

WO200075184-A1

14-DEC-2000

05-JUN-2000; 2000WO-US015449

99US-0137494P

(UYYA) UNIV YALE

Zhang

Tsvetkov

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RESULT 20
AAD12279/c
ID AAD12279/c
ID AAD12
XX AAD12
XX 25-SH
XX Human
XX H
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Modulating polypeptide levels in a cell, diagnosing and treating tumor, involves altering levels of proteins such as S-phase kinase associated proteins 1, 2 and cullin/CDC53 proteins.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; gene structure; phenotypic expression; guanosine cofactor; germline variation analysis; exon-intron boundary; Tetrahymena rRNA; p53;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 1317 BP; 295 A; 408 C; 352 G; 262 T; 0 U; 0 Other;
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                                                                                                                   primer_bind
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAD12279;
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                                            primer_bind
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/bound_moiety= "
complement(321.
                                                                                                                           640.
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/note= "Translation start
complement(137. .156)
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complement (516.
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/bound_moiety= "Primer
136. .138
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complement(615.
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bound_moiety= "Primer 6a"
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                                                                                                                                                                                                                                                                       "Primer 3a"
. .535)
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. .344)
                                               "Primer 5b" . .811)
                                                                                                                                                                                                "Primer 4a"
. .634)
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                                                                                                                                                   52=
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Determining structure of genes whose sequence is not known from cDNA, is sequencing the gene or gene across exon-intron boundaries using evenly spaced primers comprising nucleic acids that hybridize to the cDNA of
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                                                                                                                                                                                                                                             26-JUL-2001
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                                                                                                                                                                           WPI; 2001-465380/50
                                                                                                                                                                                        Thomann H,
                                                                                                                                                                                                                  20-JAN-2000; 2000US-00488127
                                                                                                                             Example
                                                                                                                                                                                                      (GENO-) GENOME THERAPEUTICS CORP
                                                                                                                                                                                                                                                           WO200153529-A2
                                                                                                                            2; Fig 2; 81pp; English
                                                                                                                                                                                        Fitzgerald
                                                                                                                                                                                                                                                                                    /bound_moiety= "Primer complement (1257. .1276)
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962. .981
                                                                                                                                                                                                                                                                                                                                                             complement (933.
/*tag= i
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The present invention relates to a method for determining gene structure CC when the genomic sequence is unknown. The method involves sequencing the Gene across exon-intron boundaries using evenly spaced primers or tiled CC primers. The tiled primers comprises nucleic acids that hybridise to the Known cDNA sequence of the gene at about 100-300 base intervals and the CC known cDNA sequence the entire gene. The method provides information the CC need to sequence the entire gene. The method provides information comprises the template. Gene structure can be determined without the need to sequence the entire gene. The method provides information without the need to sequence entire chromosomal copy of the gene or fragment. The methods are useful in germline sequence variation analysis. The method is CC acids that were separated by intervening sequence, and also for acids that were separated by intervening sequence, and also for such as Tetrahymena rRNA, where self-splicing occurs in the presence of guanosine cofactor. The present sequence is human p53 cDNA related to the

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Sequence 1317
BP; 295 A;
408 C;
352
<u>ი</u>
 262
T; 0
U; 0 Other;
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0

RESULT 21
AAD43929/c
ID AAD43
XX
AC AAD43
XX
DT 13-DI
XX 망 8 Query Match Best Local S Matches 18 AAD43929 AAD43929 127 l Similarity 18; Conserv 1 CCCGGAAGGCAGTCTGGC standard; CCCGGAAGGCAGTCTGGC Conservative DNA; 1317 100.0%; 110 18 0 쁑 Score 18; DB Pred. No. 28; Mismatches BB 4 Length 1317; 0 Gaps

13-DEC-2002

(first entry)

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RESULT 22
AAT79880/
ID AAT7
XX
AC AAT7
AC AAT7
XX
AC DT 03-D
XX
DT 03-D
XX
DE DNA
XX
BONE
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                                                                                                                                                                                                                                                                                                                                                                  The present invention relates to novel modified p53 proteins containing a composition of the present invention and a modified p53 tetramerisation domain that does not hetero-oligomerise with the native p53 tetramerisation domain. Sequences of the invention are useful for the prevention, diagnosis and treatment of diseases associated with inappropriate p53 expression and activity. They are used to treat disorders associated with decreased expression by rectifying mutations or deletions in a patient s genome that affect the activity of p53 by expressing inactive proteins or to supplement the patients own production of p53. p53 sequences are useful to prevent transplant rejection, to treat autoimmune diseases such as systemic lupus erythematosus, rheumatoid arthritis, cancers, psoriasis, atherosclerosis and arterial restenosis. They are also used as vaccines. Sequences of the invention are used in gene therapy and protein therapy.

The present sequence is human p53 wild-type DNA
                                                                                                                                                                                                                                                                                          Query Match
Best Local :
                                                                                                                                                                                                                                                                            Matches
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                                                                                                                                                                                                                                                                                                                                           Sequence 1317 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; Col 13-16; 14pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Modified p53 protein useful for preventing diagnosing and treating e.g cancers and psoriasis, comprises a modified p53 tetramerization domain that does not hetero-oligomerize with the native p53 tetramerization
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2002-462356/49.
P-PSDB; AAE26342.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Halazonetis TD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    08-MAY-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo
Bone marrow; acute myelocytic leukaemia; p53 mutant; chromosome
                                                                03-DEC-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human
                                                                                                                            AAT79880 standard; DNA; 1760 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (WIST-) WISTAR
                             DNA encoding human p53 variant found in bone marrow cell line CMC-1.
                                                                                                                                                             22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   sapiens
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                                                                                                                                                                                                                                                                           18;
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                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                            CCCGGAAGGCAGTCTGGC 110
                                                                                                                                                                                                                                          CCCGGAAGGCAGTCTGGC 18
                                                                                                                                                                                                                                                                           Conservative
                                                              (first
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136. .1317
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /product= "Human p53 wild-type protein"
                                                                                                                                                                                                                                                                                                                                       297 A; 408 C; 350 G; 262 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DNA
                                                              entry)
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                                                                                                                                                                                                                                                                                            Score 18;
Pred. No.
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S 묽

AAX75765

AAX75765 standard; DNA;

1760

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RESULT 23
AAX75765/
ID AAX7
XX
AC AAX7
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Best Local S
Matches 18
                                                                                                                                                                                                   AMT79880 is a DNA sequence encoding a p53 mutant protein. The DNA can be found in a new cell line CMC-1 (FERM p-1507), which was derived from a bone marrow cell of a patient with acute myelocytic leukaemia. The cell line has the following characteristics: (a) a chromosomal (8; 21) locus; (b) is deficient in one of 17th chromosomes; and (c) has mis-sense mutations on codons 248 and 368 of the p53 gene. The features table shows these mutations as tags c and e. Mutations shown by tags b and d are not explained in the specification. The cell line is used for production of myeloperoxidase (MPO) which has anti-fungal and anti-viral activity or as a control for detection of the (8; 21) locus in studying acute bone marrow leukemia. MPO can be supplied in a large amount as opposed to conventional production of MPO from blood as the starting material
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        JP09206070-A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                antifungal; antiviral;
                                                                                                                                                                                                                                                                                                                                                               Disclosure; Page 4-6; 6pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                   Cell line derived from leukaemic human bone marrow cells - used for the production of myeloperoxidase which has antifungal and antiviral
                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1997-460745/43.
P-PSDB; AAW25155.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  31-JAN-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   mutation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   mutation
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                                                                                                                                                                               Sequence 1760 BP; 401 A; 512 C; 461 G; 386 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                       activity.
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                                                                                    206
                                                                                                          Н
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18; Conser
                                                                                                          CCCGGAAGGCAGTCTGGC 18
                                                                                    CCCGGAAGGCAGTCTGGC 189
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                                                                                                                                  Conservative
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1031. .1033
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/note= "CGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   mutation in 
1316. .1318
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956. .958
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215. .1396
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/note= "TTY
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                                                                                                                                                                                                                                                                                                       caused by, or associated with, an RNA molecule that has a frameshift mutation. The method is used to diagnose age-related diseases, especially cancer and a wide range of neurodegenerative disorders (e.g. Alzheimer's disease, Down's syndrome, myotonic dystrophy, Huntington's disease, alcoholic liver disease, Huntington's disease, and many others listed) or susceptibility to these disorders. The method allows a definitive diagnosis of Alzheimer's disease in living patients, at an early stage. It is based on the observation that disease may be caused by mutations in RNA rather than DNA. The invention describes the used of neuronal system RNA molecules, specifically proteins including beta-amyloid precursor protein (beta-APP), the microtubule associated protein 2 (MAP2), neurofilament-I, neurofilament-M, secondated protein 2 (MAP2), neurofilament-I, neurofilament-M, cassociated protein C(MAP2), neurofilament-I, neurofilament-M, protein C(FAPP), the cellular tumour antigen p53, B-cell leukemia/lymphoma control of the cellular tumour antigen p53, B-cell neukemia/lymphoma control of the cellular tumour antigen p54 and neuromodes the wild type and mutant protein fragments represented in
                                                                                                                                                                    Query Match
                                                                                                                Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Diagnosing disease by detecting frameshift mutations in RNA or corresponding protein mutations - used to diagnose cancer and neurological diseases, particularly Alzheimer's disease, and also treatment and prevention with specific ribozymes or wild-type RNA.
                                                                                                                                                                                                                                Sequence 1760 BP; 401 A; 513 C; 460 G; 386 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This invention describes a novel method for the diagnosis of a disease caused by, or associated with, an RNA molecule that has a frameshift
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Fig 14; 258pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Van Leeuwen FW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (ROYA-)
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                                                                                                                                            Local
206
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                                                                                                                18; Conser
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                                                    CCCGGAAGGCAGTCTGGC 18
                                                                                                                Conservative
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Pred. No.
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                                                                                                                Mismatches
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                                                                                                                                                                        Length 1760;
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RESULT 25
ABA00952/c
ID ABA00
XX
AC ABA00
XX
AC ABA00
XX
DT 16-MP

standard; cDNA; 1760 BP

ABA00952; ABA00952

16-MAY-2003

(first entry)

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RESULT 24
ADD5409/c
ID AD540
XX ADD540
XX ADD540
XX ADD540
XX Human;
XW Human;
XW Colon
OS Homo s
XX Homo s
XX Homo s
XX Homo s
XX CDS
FT CDS
CC CANCE
XX CLUDW
PA (CORR
XX LUDW
PA (CO
                                                                                                             Query Match
Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                    The invention relates to a method for diagnosing colon cancer in a subject which comprises identifying colon cancer-associated polypeptides as antigens that elicit immune response in colon cancer. The method is useful for diagnosing, determining onset, progression, or regression of colon cancer in a subject, or for selecting a course of treatment of a subject having or suspected of having colon cancer. The colon cancer-associated polypeptides are useful as markers for diagnosing colon cancer, and for following the course of treatment of colon cancer. The colon cancer is human following the course of treatment of colon cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human;
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                                                                                                                                                                                                                           Sequence 1760 BP; 400 A; 513 C; 461 G; 386 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1; Page 83-84; 122pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Diagnosing colon cancer associated polypeptides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Chen Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        02-MAY-2002; 2002WO-US013994
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                                                                                                                                                                                                                                                                                  present sequence is human colon cancer-associated
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                                                      CCCGGAAGGCAGTCTGGC 18
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                                                                                                                Conservative
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gene; ds.
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/transl_except= (pos:502. .505, aa:Val)
/note= "Xaa corresponds to any amino acid"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        location/Qualifiers
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as antigens that elicit immune response in colon
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Pred.
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RESULT 26
ADD22450/c
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AC ADD224
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KW colon;
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Matches 18
                                                                                                                                                                                                                                                         comprising Sir2, a transcription factor and the compound. The method is useful for screening compounds that modulate cellular and organismal processes. Modulators of Sir2 and/or p53 activity are useful in modulating various cellular processes, e.g. repair of DNA damage, apoptosis, oncogenesis, gene silencing and senescence. Such modulators are also useful for treating cancer and aging, or for treating a cell that has been exposed to ionizing radiation
                                                                                                                                                                                                                                                                                                                                               The sequences given in ABA00951-59 encode proteins which were used method of the invention for the screening of a compound. The method comprises determining if the compound modulates silent information regulator 2 (Sir2) interaction with a transcription factor or Sir2-mediated deacetylation of a transcription factor in a reaction mixtu
                                                                                                                                                                                                                                                                                                                                                                                                                                       Screening for modulators of silent information regulator 2 (Sir2) protein for treating cancer and aging, comprises determining if the compound modulates Sir2-mediated deacyetylation of the transcription factor.
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06-JUL-2001; 2001US-0303456P
05-JUL-2002; 2002US-00303456
tumour antigenic peptide; cancer; vaccine; cytostatic; cytotoxic T cell; colon; mouth; lung; prostatic; gynecological; human; gene; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Fig 14; 93pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2003-210354/20
P-PSDB; AAG79917.
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                                HLA-B46 T cell recognised tumour antigenic polypeptide, SEQ No 100
                                                        15-JAN-2004
                                                                                                                                                                                                                                               Sequence 1760 BP;
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                                                                                                   ADD22450 standard; DNA; 2451
                                                                               ADD22450,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          screening; silent information regulator 2; Sir2; deacetylation; cription factor; modulator; p53; DNA repair; apoptosis; oncogenesis; silencing; senescence; cancer; aging; ionizing radiation; ss.
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215. .1396
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The invention further provides a cancer vaccine comprising a tumour CC antigenic peptide or polypeptide, which has cyrostatic activity. A tumour CC antigenic peptide, polypeptide, its encoding polynucleotide, a ccompound of the tumour CC polynucleotide, a recombinant vector containing the CC polynucleotide, a recombinant vector or an antibody are CC useful for screening for compounds that interact with the tumour antigenic peptide, the polypeptide or its encoding polynucleotide and CC increases the expression of the tumour antigenic peptide, the polypeptide or polynucleotide. The tumour antigenic peptide or polypeptide is useful for inducing cytotoxic T cells. The tumour antigenic peptide is useful for inducing cytotoxic T cells. The tumour antigenic peptide or prostatic or gynecological cancer. The invention also provides a composition useful for treating cancer. The tumour cantigenic peptide or the polypeptide is useful as an antigen to create antigenic peptide or the polypeptide is useful as an antigen to create cantibodies. This polynucleotide sequence represents the DNA encoding one of the human tumour antigenic polypeptides of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Novel tumor antigenic peptide or polypeptide useful for induci: cytotoxic T cells or for treating cancer such as colon, mouth, prostatic or gynecological cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to a novel tumour antigenic peptide or polypeptide comprising a sequence selected from 99 sequences fully defined in the specification. The tumour antigenic peptide or polypeptide comprises a sequence selected from 99 sequences fully defined in the specification, where the tumour antigenic peptide preferably has a sequence of Glu-ro-ro-leu-ser-Gln-Glu-Thr-Phe, and the polypeptide preferably has a sequence comprising 393 amino acids fully defined in the specification.
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Sequence
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2451 BP; 537 A; 697 C; 592 G; 625 T; 0 U; 0 Other;
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Ş Query Match Best Local S Matches 18 16; ы Similarity CCCGGAAGGCAGTCTGGC 18 Conservative 100.0%; 0 Score 18; Pred. No. Mismatches 9; 0 Length 2451; 0 Gaps

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                                                                                    DNA encoding novel human diagnostic protein #1503
                                                                                                            13-FEB-2002
                                                                                                                                 AAS65699;
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Human; chromosome mapping; gene mapping; gene therapy; forensic; food supplement; medical imaging; diagnostic; genetic disorder; forensic; 88

WO200175067-A2

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RESULT 28
AAS67409/c
ID AAS67
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XX AAS67
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XX Humar
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P-PSDB;
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23-AUG-2000;
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                                                                                                                                                                 Homo sapiens
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DB; ABG01512.
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supplement; n
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18; Conservative
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2000US-00649167.
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                                                                                                                                                                                                                                                                                                                               novel human diagnostic protein #3213
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imaging; diagnostic;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local S
Matches 18
Human; cytokine; cell proliferation; cell differentiation; growth factor; haematopoiesis regulation; tissue growth; immunomodulator; activin; inhibin; chemotaxis; chemokinesis; thrombolysis; oncogenesis; proliferation; metastasis; cancer; tunour; haematopoietic disorder; proliferation; metastasis; cancer; tunour; haematopoietic disorder; chronic inflammatory condition; proliferative retinopathy; atherosclerosis; coronary heart disease; arterial ischaemia; bone disorder; osteoporosis; vascular growth disorder; tissue regeneration; wound healing; infection; immune disorder; cell culture; drug screening; gene therapy; antiinflammatory; antiasthmatic; antiarthritic; haemostatic; antiarteriosclerotic;
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23-AUG-2000; 2000US-00649167
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                                                                                                                                                                                                                                                                                                                                                                                                                        Human p53 homologue-encoding cDNA, SEQ ID NO:1155.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABA09379;
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Pred. No.
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cytostatic; osteopathic; vasotropic; cardiant; virucide; antibacterial; antifungal; vulnerary; antiulcer; ss.

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Sequences ABB10981-ABB12330 represent musleic acids encoding them. The civention also relates to vectors and recombinant host cells comprising a cor nucleotide of the invention, methods of producing the novel polypeptides, antibodies against the polypeptides, methods of detecting the novel polypeptides or polypeptides in a sample, and methods of identifying compounds which to be polypeptides of the invention Although novel, many of the polypeptides of the invention Although novel, many of the polypeptides of the invention Although novel, many of the polypeptides of the invention Although novel, many of the polypeptides of the invention are homology to known proteins, thereby complete a regulatory activity; is such as the polypeptides of the invention may conditions. The polypeptides of the invention may conditions or cell differentiation activities, including cytokine, cell proliferation or cell differentiation activities, including cytokine, cell proliferation or cell conditions regulatory activity; tissue growth factor activities, themeotopies cancivities, including cytokine, cell proliferation or cell involved in oncogenesis, cancer cell proliferation, thrombotide or thrombotylic activities, receptor or ligand activities, or may be personably representing treating or materials.

Compending on their biological activities, haemostatic, thrombotides of the invention are useful for preventing, treating or ameliorating medical conditions, e.g., by protein or gene therapy. Such conditions include cancers, haematopoietic disorders (e.g., myeloid or lymphoid cell disorders, haematopoietic activities, polypeptides and nucleatides of conditions, e.g., by protein or gene therapy. Such conditions and abnormal vascular growth. Polypeptides involved with tissue regeneration and creative retinopathy, atheracolerosis, coronary hear disease.

Compending the conditions of the compending the polypeptides and nucleatives to polypeptides with growth factor activity may be used to promote wound to many polypeptides and nucleative
                  Query Match
Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human proteins and DNA encoding sequences useful for preventing, treating or ameliorating a medical condition in a mammalian subject e.g. arthritis
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P-PSDB; ABB12135.
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27-APR-2000; 2000US-00560875.
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1 Similarity
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Score 16.4; D
Pred. No. 1.6e
0; Mismatches
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                                                                                                                                                                         ) STACHE-CRAIN B.
) DICKSON M C.
) JONES L W.
                                                                                                                                                                                          DRMANAC R T.
LABAT I.
                                                                                                                                                                                                                                                                                                                    standard;
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                                                                                                                                                               Dickson MC,
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                                                                                                                                                                                                                                                                        EST,
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New polynucleotide sequences obtained from various cDNA libraries, useful as hybridization probes, as oligomers for PCR, for chromosome and gene mapping, in the recombinant production of protein, or in generating antisense DNA or RNA.

Claim 1; SEQ ID NO 36880; 44pp; English

The invention relates to an isolated polynucleotide comprising any one of CC 38043 cDNA sequences, appearing as ACHI-2789-ACH50831, whose sequence was CC determined by the technique of SBH (sequencing by hybridisation). Also concluded is a purified polypeptide comprising a sequence corresponding to a reading frame of the novel polynucleotide. The nucleic acid sequences are useful in diagnostics as expressed sequence tags (EST) for comprising expressed genes or for physical mapping of the human genome, in forensics, in assessing biodiversities, or in identifying mutations cresponsible for genetic disorders and other traits. The nucleotide sequences are also useful as hybridisation probes, as oligomers for PCR, for chromosome and gene mapping, in the recombinant production of protein, or in generating antibodies specific for it. The present sequence is useful for generating antibodies specific for it. The present sequence is one of the 38043 isolated cDNA/EST sequences. Note: The sequence obtained in electronic format directly from USPTO at sequence.html?DocID=20030073623

Sequence 436 BP; 84 A; 148 C; 113 <u>.</u> 91 T; 0 U; 0 Other;

맑 á 90 CCCGGAAGGCAGTCTGGC

Matches Query Match Best Local

Similarity

91.1%; 94.4%;

Score 16.4; DB 8; Pred. No. 1.6e+02;

Length 436; Indels

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Conservative

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Human; connec cytostatic; 9 Homo sapiens: WC200155343-J 02-AUG-2000; 04-FEB-2000; 04-FEB-2000; 04-FEB-2000; 04-FEB-2000; 04-FEB-2000; 04-FEB-2000; 04-FEB-2000; 04-FEB-2000; 04-FEB-2000; 04-FEB-2000; 04-FEB-2000; 01-MAR-2000; 11-MAR-2000; 11-MAR-2000; 11-MAR-2000; 07-JUL-2000; 07-JUL-2000; 07-JUL-2000; 07-JUL-2000; 07-JUL-2000; 07-JUL-2000; 07-JUL-2000; 07-JUL-2000; 07-JUL-2000; 07-JUL-2000; 07-JUL-2000; 07-JUL-2000; 01-AUG-2000; 01-AUG-2000; 14-AUG-2000; 14-AUG-2000; 14-AUG-2000; 14-AUG-2000; 14-AUG-2000; 14-AUG-2000; 14-AUG-2000; 14-AUG-2000; 14-AUG-2000; 14-AUG-2000; 14-AUG-2000; 14-AUG-2000; 14-AUG-2000; 14-AUG-2000; 14-AUG-2000; 01-SEP-2000; 01-SEP-2000; 01-SEP-2000; 01-SEP-2000; 06-SEP-2000; 08-SEP-2000;	ABK42958/c ID ABK42958 star XX AC ABK42958; XX OT 21-MAY-2002 XX
ic gene; ds. iens. 343-A1. 001. 001. 2000US-0179065P, 000. 2000US-0188354P, 000. 2000US-018835P, 000. 2000US-0205515P, 000. 2000US-021886P, 000. 2000US-021889P, 000. 2000US-022851P, 000. 2000US-022851P, 000. 2000US-022858P, 000. 2000US-02898P, 000. 2000US-022898P, 000. 2000US-028898P, 000.	standard; DNA; 942 BP. 02 (first entry) equence #857 encoding novel human connective
e thezapy;	tissue polypeptide.
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25-SEP-2000; 26-SEP-2000; 27-SEP-2000; 29-SEP-2000; 29-SEP-2000; 29-SEP-2000; 29-SEP-2000; 29-SEP-2000; 20-CCT-2000; 20-CC	
2000US-023436PP 2000US-0235834PP 2000US-0235836PP 2000US-023636PP 2000US-023636PP 2000US-02363703PP 2000US-02363703PP 2000US-0237039PP 2000US-023935PP 2000US-023935PP 2000US-023935PP 2000US-0231786PP 2000US-0231786PP 2000US-0241786PP 2000US-0241786PP 2000US-0241826PP 2000US-0241826PP 2000US-0246477PP 2000US-0246477PP 2000US-0246477PP 2000US-024661PP 2000US-024652PP 2000US-024921PP 2000US-024929PP 2000US-024929PP 2000US-024929PP	000000000000000000000000000000000000000

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RESULT 32
ABK42959/c
ID ABK42959;
XX
AC ABK42959;
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AC ABK42959;
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DT 21-MAY-200
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Matches
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05-DEC-2000;
06-DEC-2000;
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  31-JAN-2000;
04-FEB-2000;
24-FEB-2000;
02-MAR-2000;
16-MAR-2000;
17-MAR-2000;
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used in
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                                                                                                                                                                                                                                                                                                   Human; connective tissue related disorder; cancer; gene therapy; cytostatic; gene; ds.
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2000US-0251719P.
2000US-0251479P.
2000US-0251856P.
2000US-0251868P.
2000US-0251869P.
2000US-0251869P.
2000US-0251989P.
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2000US-0254097P.
2001US-0259678P.
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2000US-0179065P.
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2000US-0186350P.
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                                                                                                                                             2001WO-US001322
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94.4%;
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a disorder such
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used in
The present invention relates to the isolation of novel human connective tissue related polypeptides (AAU86435-AAU86923) and the polynucleotide (CDNA and genomic) sequences encoding them. The sequences of the
                                                                 Disclosure;
                                                                                                  cancer
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diagnosing, preventing, tr
ar or rheumatoid arthritis.
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RESULT 33
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ID ADB61115;
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XX Cytostatic
XX Cytostatic
XX Cytostatic
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cytostatic; neuroprotective; nootropic; antiparkinsonian; cardiovascular;
KW antiarteriosclerotic; immunosuppressive; antirheumatic; antiarthritic;
KW antiinflammatory; antiallergic; antiasthmatic; dermatological;
KW nephrotopic; virucide; fungicide; antibacterial; antiparasitic;
KW gene therapy; ds; connective tissues disorder; heumatoid arthritis;
KW gene therapy; ds; connective tissues disorder; heumatoid arthritis;
KW cancer metastasis; neoplasia; leukaemia; neurodegenerative disorder;
KW Alzheimer's disease; parkinson's disease; cardiovascular disease;
KW Alzheimer's disease; parkinson's disease; cardiovascular disease;
KW autcimmune disease; multiple sclerosis; allergic reaction; asthma;
KW autcimmune disease; multiple sclerosis; allergic reaction; asthma;
KW rhinitis; eczema; inflammatory condition; Crohn's disease; nephritis;
KW gastrointestinal disorder; inflammatory bowel disease;
KW vogan transplant rejection; immune system disorder; Bruton's disease;
KW X-linked lymphoproliferative syndrome;
W Chromosome identification; chromosome mapping;
KW connective tissue related polynucleotide; gene; ds.
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Matches 17
  18-APR-2000
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17-MAR-2000;
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2000US-0184664P.
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2000US-0186350P.
2000US-0199874P.
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2000US-0209467P.
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17-NOV-2000
The invention describes an isolated nucleic acid molecule (I), which comprises a sequence that is at least 95 % identical to a connective tissue-related polynucleotide encoding connective tissue antigens (CTA) The polypeptide or polynucleotide is useful for preventing, treating, ameliorating medical conditions in a mammal. The connective tissue polypeptides, polynucleotides and antibodies are particularly useful for treating, preventing and/or prognosing disorders of connective tissues (e.g. rheumatoid arthritis, discoid and systemic lugus erythematosus, scleroderma, or Sjogren's syndrome), cancers, cancer metastases and/or neoplasias (e.g. leukaemia), neurodegenerative disorders (e.g. halzheimer's disease, or Parkinson's disease), cardiovascular diseases
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17-NOV-2000;
17-NOV-2000;
                                                                                                                                                                                                                                                                                                                                                                           New connective tissue-related polypeptides and polynucleotides, useful for treating, preventing and/or prognosing e.g. disorders of connective tissue, (e.g. rheumatoid arthritis), cancers, cancer metastases and/or
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P-PSDB; ADB59710.
                                                                                                                                                                                                                                                                                                         Disclosure;
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13-AUG-2000;
14-SEP-2000;
16-SEP-2000;
17-SEP-2000;
18-SEP-2000;
19-SEP-2000;
11-SEP-2000;
11-SE

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RESULT 34
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XW Connec
XW Cytost
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14-JUG-2000

14-JUG-2000
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2000US-0225270P.
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17-NOV-2000;
17-NOV-2000;
                                                                                     The invention describes an isolated nucleic acid molecule (I), which comprises a sequence that is at least 95 % identical to a connective tissue-related polynucleotide encoding connective tissue antigens (CTA). The polypeptide or polynucleotide is useful for preventing, treating, or ameliorating medical conditions in a mammal. The connective tissue polypeptides, polynucleotides and antibodies are particularly useful for treating, preventing and/or prognosing disorders of connective tissues (e.g. rheumatoid arthritis, discoid and systemic lugus erythematosus, scleroderma, or Sjogren's syndrome), cancers, cancer metastases and/or neoplasias (e.g. leukaemia), neurodegenerative disorders (e.g. Alzheimer's disease, or parkinson's disease), cardiovascular diseases
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17-NOV-2000;
17-NOV-2000;
                                                 (e.g. atherosclerosis, myocarditis or cardiopulmonary bypass complications), autoimmune diseases (e.g. systemic lupus erythematosus rheumatoid arthritis, or multiple sclerosis), allergic reactions (e.g.
                                                                                                                                                                                                                                                         Disclosure;
                                                                                                                                                                                                                                                                                             New connective tissue-related polypeptides and polynucleotides, useful for treating, preventing and/or prognosing e.g. disorders of connective tissue, (e.g. rheumatoid arthritis), cancers, cancer metastases and/or
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17-NOV-2000;
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2000US-0246528P.
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2000US-0254097P
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           Score 16.4; DB 8;
Pred. No. 1.7e+02;
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Matches

Conservative

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Indels

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Gaps

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30-JUN-1995 AAQ72979;

(first entry)

AAQ72979

standard; cDNA; 140

ВP

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RESULT 35
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ID AAH230
AC Osteoa
AC AAH230
AC AA
RESULT 36
AAQ72979
ID AAQ72
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AC AAQ72
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Substantially purified protein, polypeptide or their fragments, identify a biologically active compound or composition and treat mammalian osteoarthritis.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 1986
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RESULT 37
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Matches 16
                                                                                                                                                                                                                                                                                                                    AAQ72979 which encodes AAR62645 is the antisense strand to a p53 gene human tumour antigen. In healthy women expression of AAR62645 is repressed by the normal wild type p53 gene, however mutations in this gene could permit the expression of this antisense protein, which may lead to the development of breast cancer. Therefore AAR62645 can be used in the development of humanised antibodies agianst breast cancer, which due to AAR62645's non-expression in healthy women, should have no side
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                                            misc_difference
                                                                       Homo sapiens
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16; Conser
                                                                                                                                                                                                                                                                                            140 BP; 30 A; 46 C; 44 G; 20 T; 0 U;
                                                                                                                                                                                                                                      CGGAAGGCAGTCTGGC 18
                                                                                                                                                                                                                      CGGAAGGCAGTCTGGC 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          tumour antigen antisense cDNA
                                                                                                                                                                                                                                                         ilarity 100.0%; I Conservative 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Bergmann
                                                                                                                            (revised)
(first entry)
                                                                                         BC538.1;
                                                                                                         strand cDNA encoding protein BC538.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            93CA-02090879
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              93CA-02090879
                                            Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Location/Qualifiers
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transl_except= Val
                                                                                                                                                                          CDNA; 140
                                                                                                                                                                                                                                                                          88.9%;
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                                                                                         p53 antisense strand; breast
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       antisense breast cancer therapy;
                                                                                                                                                                                                                                                                  Score 16; Pred. No
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                                                                                                                                                                                                                                                          Mismatches
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                                                                                                                                                                                                                                                        2.5e+02;
hes 0;
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Matches
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                                               17-DEC-1998;
17-DEC-1998;
23-JUN-1999;
24-SEP-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAQ74243, AAQ74244 and AAQ74245 are cDNA fragments of the p53 gene antisense strand which encode AAR51625 (BC534), AAR51626 (BC538) and AAR51627 (BC534), DE538 (BC538) and AAR51627 (BC538 I) respectively. These proteins can be used to develop products which can be used in the diagnosis and prediction of breast cancer and other cancers associated with mutatations of the p53 gene. These cancers can be treated by providing the patient with an inhibitor of the BC534, BC538 and BC538.1 genes, and the regulatory sequence AAQ74246 (BC53/reg). (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New p53 anti sense proteins - used to develop prods. for the diagnosis, prediction and treatment of breast cancer and related cancers.
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(PRED/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            misc_difference
                                                                                                                                                                 22-JUN-2000.
                                                                                                                                                                                                 WO200036107-A2
                                                                                                                                                                                                                                 Homo sapiens.
                                                                                                                                                                                                                                                                                 Human;
                                                                                                                                                                                                                                                                                                                   Human
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                                                                                                                                                                                                                                                                                                                                                                                   AAA70021;
                                                                                                                                                                                                                                                                                                                                                                                                                   AAA70021 standard; cDNA; 184
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 140 BP; 30 A; 46 C; 44 G;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Bergmann JE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            16-MAR-1993;
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                                                                                                                                 17-DEC-1999;
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                                                                                                                                                                                                                                                                                                                   ovarian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       16 CGGAAGGCAGTCTGGC 31
                                                                                                                                                                                                                                                               ovarian carcinoma; ovarian cancer; therapy; diagnosis;
antigen; identification; cytostatic; gene therapy; vaccine;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     l Similarity
16; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CGGAAGGCAGTCTGGC 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ilarity 100.0%;
Conservative
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                                              98US-00215681.
98US-00216003.
99US-00338933.
99US-00404879.
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100.0%; Pred. No. 2.1
tive 0; Mismatches
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hes 0;
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CORIXA

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RESULT 39
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Best Local S
Matches 16
                                                                                                                                                                                                                                                                                    Mitcham
Reed SG,
                                                                                                                                                                                                                                                                                                                                                                                             17-JUL-2000; 2000US-00617747, 10-AUG-2000; 2000US-00658601. 2005E9-000; 2000US-00667857, 04-APR-2001; 2001US-00887271, 18-JUN-2001; 2001US-00884441
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention describes an isolated polypeptide comprising an immunogenic portion of an ovarian carcinoma protein (or its variants). Ovarian carcinoma proteins, and polynucleotides encoding them, have cytostatic activity and can be used in gene therapy and vaccines. Ovarian carcinoma polypeptides, nucleic acids, antibodies and vaccines are useful for the prevention, diagnosis and treatment of cancer, preferably ovarian cancer. AAA69691 to AAA70077 and AAB12552 to AAB12557 represent human ovarian carcinoma polynucleotides and proteins used in the exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Immunogenic portion of an ovarian carc encoding it, useful for the diagnosis, cancer, preferably ovarian cancer.
                                                                                                                                      Polypeptides comprising an immunogenic portion of protein or its variants, useful for stimulating patient and treating ovarian cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           17-JUL-2001; 2001WO-US022635.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     24-JAN-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; immunostimulant; cytostatic; cancer; ovarian carcinoma;
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                                                                                            Example 2; Page 298; 408pp; English
                                                                                                                                                                                                                                     WPI; 2002-164781/21.
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, Vedvick
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP; 61 A; 27 C;
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                                                                                                                                                                                                                                                                                    is,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         antigen polynucleotide #20.
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94.1%;
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Pred. No. 5e
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                                                                                                                                                                                                                                                                                  Fling
Hill P,
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5e+02;
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                                                                                                                                                                                                                                                                                  SP, Retter
, Albone E;
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                                                                                                                                                               an ovarian carcinoma
n immune response in
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                                                                                                                                                                                                                                                                                                      Fanger
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This invention relates to polypeptides comprising an immunogenic portion of an ovarian carcinoma protein which acts as an immunostimulant and is cytostatic. The polypeptides, polynucleotides, antibodies, fusion

Sequence 184

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Best Local
                                                          The invention relates to an isolated 0772P polypeptide, which has the structure fully defined in the specification. The composition containing the polypeptides, polynucleotides, fusion proteins, antibodies T-cells or antigen presenting cells are useful for stimulating an immune response and treating ovarian cancer. Detecting the presence of the polynucleotides and polypeptides is useful for diagnosing cancer. Ovarian carcinoma cDNAs and protein cDNAs were identified using microarray technology. The present sequence represents a human ovarian carcinoma
                                                                                                                                                                                                                                                                                                                                                                                                                                             Mitcham
Reed SG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        proteins, T cell populations and antigen presenting cells that express the polypeptides are useful for stimulating an immune response in a patient and treating ovarian cancer. This sequence represents DNA related to the invention
                                                                                                                                                                                                                                                                                                                            New isolated 0772P polypeptides and polynucleotides, therapy, particularly for treating or diagnosing can
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    06-NOV-2003
                                                                                                                                                                                                                                                               Example 2; SEQ ID NO 332; 371pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ds; human; gene therapy; ovarian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human
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3, Vedvick
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KING G E.
ALGATE P A.
FLING S P.
FLING S P.
FANGER G R.
FANGER G R.
REED S G.
VEDVICK T S.
CCARTER D.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          HILL P.
ALBONE E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CCGGAAGGCAGTCTGGC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2001US-00884441
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                                                                                                                                                                                                                                                                                                                                                                                                                                             IS GE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A; 27 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               85.6%;
94.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                             Algate
Carter
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              47 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 15.4; DB | Pred. No. 5e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                           DΑ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                             Fling :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              49
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                                                                                                                                                                                                                                                                                                                                                                                                                        j SP,
b. Albone
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                                                                                                                                                                                                                                                                                                                                                                                                                                           Retter
bone E;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Other;
                                                                                                                                                                                                                                                                                                                                    cancer,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Fanger
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DB 8;

Length 184;

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RESULT 41
AA129201/c
ID AA1292
XX AA1292
XX AA1292
XX AA1292
XX Colon
XX Human;
XW Gene t
XX WO2001
XX WO2001
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XX WO2001
XX WO2001
XX WO2001
XX AO2001
XX AO2
                  The present invention describes colon tumour associated proteins (I) and (II) can be used in gene therapy and vaccine production. (I) and (II) may be used in the prevention, diagnosts and treatment of diseases (II) may be used in the prevention, diagnosts and treatment of diseases (C associated with inappropriate colon tumour associated protein (TCAP) (C expression, such as colonic cancer. For example, (I) and (II) may be used to treat disorders associated with decreased expression by rectifying (C mutations or deletions in a patient's genome that affect the activity of (CTAPs) by expressing inactive proteins or to supplement the patients own production of them. Additionally, (II) may be used to produce the TCAP (C proteins, by inserting the nucleic acids into a host cell culturing the C cell to express the protein. (II) and its complementary sequences may (C and hybridisation assays to detect and quantitate the presence of similar (C nucleic acids in samples, and therefore which patients may be in need of c restorative therapy, (I) may also be used as antigens in the production of antibodies against TCAPs and in assays to identify modulators of TCAP (C antibodies may also be used as antibodies and antagonists may also be used to down regulate TCAP expression and activity. Anti-(I) antibodies and antagonists may also be (EIISA), AA128460 to AA129512 and AAV4494 to AAV44233 represent assay (I presence of TCAPs in samples (e.g. by enzyme linked immunosorbant assay (I presence of TCAPs in samples (e.g. by enzyme linked immunosorbant assay (I presence of TCAPs).
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Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           10-JAN-2000;
15-FEB-2000;
06-MAR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 2; Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Colon tumor associated proteins and nucleic acids useful prevention, diagnosis and treatment of colonic cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAI29201
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAI29201;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               gene therapy;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Lodes MJ, surprise T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               immunotherapy; diagnosis; colon cancer;
herapy; vaccine; colonic cancer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORIXA CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            standard; cDNA; 405
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CCAGAAGGCAGTCTGGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; 2000US-00480321.
2000US-00504629.
2000US-00519444.
2000US-00575251.
2000US-00609448.
2000US-00649811.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       347; 472pp; English.
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, Jiang Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        85.6%;
acid sequences
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Benson DR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
given
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Meagher MJ,
in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            colon tumour; immunogenic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
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RESULT 42
ABZ33387/c
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                                                              Query Match
Best Local (
                                                 Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                      diagnosis of colon cancer. Also described: (1) a method for detecting the presence of cancer in a patient; (2) a method for stimulating and/or expanding T cells specific for a tumour protein; (3) and the T cell population comprising T cells prepared by the method of (2); (4) a method for stimulating an immune response in a patient; (5) a method for treating cancer in a patient; and (6) a method for inhibiting the development of cancer in a patient. (1) have immunostimulant and cytostatic activities and can be used in vaccines. AB332646 to AB233725 and ABP55343 to ABP55391 represent human colon cancer/tumour related sequences used in the exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                        Xu J, ;
Wang T,
Fanger (
                                                                                                                                                                                                                                                                                                         New polynucleotide, useful for the preparation of stimulating an immune response against, or treating
                                                                                                                                                                                                                                                                                                                                                                                                                                                   10-APR-2001; 2001US-00833263
03-AUG-2001; 2001US-00922217
19-DEC-2001; 2001US-00025380
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 405 BP; 106 A; 100 C; 77 G; 117 T; 0 U; 5 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      present invention
                                                                                                   Sequence
                                                                                                                                                                                                                                                                                 Disclosure; Page 349; 537pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       09-APR-2002; 2002WO-US011475
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human colon tumour cDNA for clone R0093:H07 SEQ ID NO:755
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABZ33387
                                                                                                                                                                                                                                                        The present invention describes compounds (I) for the immunotherapy
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                                                                                                                                                                                                                                                                                                                                                                                                                           (CORI-)
213
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                                                                                                                                                                                                                                                                                                                                                                        Lodes MJ, Se
I, Jiang Y, S
r GR, Vedvick
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          colon cancer; colon tumour; immunotherapy; diagnosis; cancer;
; immune response; immunostimulant; cytostatic; vaccine; gene;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    N
                                                 l Similarity
16; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                            CORIXA CORP.
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                                                                                                   405
CCAGAAGGCAGTCTGGC 197
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                        CCGGAAGGCAGTCTGGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           standard; cDNA; 405
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                                                 Conservative
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                                                                                                   106
                                                                                                                                                                                                                                                                                                                                                                        Secrist H, Benson DR, Me
Smith CL, King GE, Wang
.ck TS, Carter D;
                                                            85.6%;
94.1%;
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                                                                                                  A; 100 C; 77
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                                                             Score 15.4; DB 7;
Pred. No. 5.1e+02;
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Pred. No. 5.
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                                               Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                   Meagher MJ, Stolk JA;
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                                                                                                                                                                                                                                                                                                           treating, cancer.
                                                                        Length 405;
                                                                                                   5 Other
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CC nuclectide sequence from a genome of an organism corresponds to a nucleotide sequence from a genome of an organism corresponds to a nucleotide sequence of an open reading frame; for preparing a contig, call or part of a genome of an organism. mRNA is obtained from mammalian or human cell which is associated with a pathological condition e.g. a colon cancer or breast cancer cell. The method is useful for analyses of colon cancer or breast cancer cell. The method is useful for analyses of colon cancer or subjects and can be used to carry out genetic analyses of compatitions of subjects and can be used to study living systems to determine if, e.g. there have been genetic shifts which render conditional or population more or less likely to be afflicted with collegases such as cancer, to determine antibiotic resistance or non-conditions are likely to be passed to office office congenital diseases, and the risk of affliction to a foetus, as well as the study of whether the conditions are likely to be passed to offspring through ova or sperm. The analyses for pathological conditions can be carried out in all animals, plants, birds, fish, etc. Using this method, in the area of agriculture, for example the genomes of food crops can be studied to determine if resistance genes are present, defects in plant conditions can be studied to determine if resistance senses are breached, the method permits determination of the pathogens which integrate into the genome, such as created and other integrating viruses such as influenza virus, have conditioned should be also be studied in this way. Similarly, the method permits have undergone shifts or mutations, which may require different approaches to therapy. This method is also applied to eukaryotic pathogens, such as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention describes a method of determining open reading frames in the genome of organism, comprising contacting mRNA from cell of organism with a single oligonucleotide primer (I) at low stringency, preparing single-stranded cDNA by reverse transcribing mRNA with (I), amplifying cDNA, sequencing the product, and repeating the contacting, preparing and amplifying steps with different primers and sequencing resulting nucleic acids. The method is useful for determining that a known nucleic acids. The method is useful for determining that a known
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     breast cancer; population genome analysis; genetic shift; cancer; antibiotic resistance; antibiotic non-tolerance; congenital disease; agriculture; food crop genome; resistance gene; retrovirus; influenza virus; eukaryotic pathogen detection; trypanosome; Plasmodium;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Determining open reading frames of genome of an organism e.g. a human suffering from cancer involves use of single oligonuclectide primer at low stringency for preparing single-stranded cDNA from mRNA of individual.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Simpson
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NETO E D.
BRENTANI R R.
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RESULT 44
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Best Local S
Matches 16
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29-AUG-2000;
08-SEP-2000;
26-SEP-2000;
06-OCT-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          trypanosomes, different types of Plasmodium, etc. The method essemeliminates sequencing of non-coding portions. This sequence represpolynucleotide isolated from human colon cancer cell cDNA library
                                  The invention describes an isolated polynucleotide and polypeptide useful for stimulating and/or expanding T cells specific for a tumour protein for determining the presence of a cancer in a patient. A composition containing the polynucleotide and/or polypeptide is useful for treating a lung cancer in a patient. The polypeptide is useful for removing tumour cells from a biological sample. The polypeptide is also useful as probe or primer to detect the level of mRNA encoding a tumour protein. This sequence encodes a lung tumour associated protein or protein fragment, described in the method of the invention. Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at
                                                                                                                                                                                                                    Novel polynucleotide encoding a lung stimulating and/or expanding T cells
                                                                                                                                                                                                                                                                                                  Wang T, Wata
Marnerakis M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 428 BP; 123 A; 101 C; 79
                                                                                                                                                                                                                                                              WPI; 2002-164634/21.
                                                                                                                                                                                                                                                                                       Wang
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16; Conserv
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Fanger N, Sv
                                                                                                                                                                                                                                                                                                               Watanabe Y,
                         .int/pub/published_pct_sequences
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2000US-00671325.

2000US-00677419.

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2000US-00736457.
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                                                                                                                                                                                            ID NO 1634; 223pp; English.
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2000US-00651563
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94.1%;
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D, Fanger GR,
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Pred. No. 5
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Mcneill PD, Clapp
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5.2e+02;
tes 1;
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ck TS, Bangur CS;
Clapper JD;
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RESULT 45
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Best Local S
Matches 16
The invention relates to a polynucleotide encoding a lung tumour protein, comprising a sequence selected from any of the 14 sequences mentioned in the specification, or a sequence (S2) mentioned in specification, or a sequence (S2) mentioned in specification, or degenerate variants of at least 20 contiguous residues of S1, sequences that hybridise to S1, sequences having 75%, preferably 90%, identity to S1, or degenerate variants of S1. Also included are an isolated polypeptide (comprising a sequence (S3) selected from any one of the 4 amino acid sequences mentioned in the specification, a sequence encoded by the polynucleotide, or sequences having at least 70%, or preferably 90%, identity to a sequence encoded by the polynucleotide), an expression control sequence, a host cell transformed or transfected with the vector, an isolated antibody (or its antigen-binding fragment) that specifically binds to the polypeptide, and the vector, an isolated antibody (or its antigen-binding fragment) that specifically binds to the polypeptide, and conditions, stimulating and/or expanding T cells specific for a tumour conditions, stimulating and/or expanding T cells specific for a tumour to parmit the stimulation and/or expanding of T cells) and inhibiting the potention or antigen-presenting cells, under conditions and inhibiting the conditions and for a time sufficient to permit the stimulation and/or expanding T cells and inhibiting the cells isolated from a patient with the polynucleotide, protein or antigen to a cancer in a patient with the polynucleotide, protein or cells isolated from a patient with the polynucleotide, protein or antigen to the polynucleotide from a patient with the polynucleotide, protein companies to the polynucleotide from a patient with the polynucleotide from antigen to the cells isolated from a patient with the polynucleotide, protein or antigen to the polynucleotide from a patient with the polynucleotide from antigen to the polynucleotide from a patient with the polynucleotide from antigen to th
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Clapper JD;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel lung cancer polynucleotide encoding lung cancer protein, useful for detecting the presence of lung cancer in a patient, and in pharmaceutical compositions, e.g. vaccines, for treating lung cancer.
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(FANG/)
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(WANG/)
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16; Conservative
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WANG T.
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94.1%;
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Pred. No. 5.2e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CC presenting cells that express the polynucleotide, such that T cells croliferate, administering to the patient an effective amount of the proliferated T cells, and thus inhibiting the development of a cancer in the patient. The polynucleotide, protein ands cells are useful in a composition for stimulating an immune response in a patient, and for creating a cancer in a patient (particularly lung cancer). The coligonucleotide is useful for determining the presence of a cancer in a patient. The protein and oligonucleotides are useful in parmaceutical compositions, e.g. vaccines. The polynucleotide is also useful as a probe or primer for nucleic acid hybridisation, and in the design and corpoparation of ribozyme molecules for inhibiting expression of tumour colls. The polynucleotide is useful for isolating a full-length gene from a suitable collbrary. The present sequence is a cDNA (full length, extended or collbrary. The speech term a library derived from lung tumour/cancer cells. Once: The sequence data for this patent did not form part of the printed concerning the present sequence in electronic format directly from the USPFO at sequence as obtained in electronic format directly from the
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08-SEP-2000;
26-SEP-2000;
06-OCT-2000;
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                                                                                                                                                                                                                                                                                                                                                                     27-APR-2000;
05-JUN-2000;
11-JUL-2000;
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15-OCT-1999;
17-DEC-1999;
30-DEC-1999;
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22-MAR-2000; 2000US-00533077.
10-APR-2000; 2000US-00546259.
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16; Conservative
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2000US-00589184.

2000US-00614124.

2000US-00651563.

2000US-00658824.

2000US-006771325.

2000US-00677419.

2000US-00736457.
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Henderson Durham M,

RA, Wang T, Watanabe Carter D, Fanger GR,

Watanabe Y, Johnson JC, Retter anger GR, Vedvick TS, Bangur CS,

Retter MW;

Mcnabb

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The invention relates to an isolated polynucleotide comprising any one of 38043 cDNA sequences, appearing as ACH12789-ACH50831, whose sequence was determined by the technique of SBH (sequencing by hybridisation). Also included is a purified polypeptide comprising a sequence corresponding a reading frame of the novel polynucletide. The nucleic acid sequences are useful in diagnostics as expressed sequence tags (EST) for identifying expressed genes or for physical mapping of the human genome, in forensics, in assessing biodiversities, or in identifying mutations
                                                                                                                                                                                                                                                                                                            New polynucleotide sequences obtained from various as hybridization probes, as oligomers for PCR, for mapping, in the recombinant production of protein,
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16; Conserv
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mapping; biodiversity; genetic disorder.
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94.1%;
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Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         responsible for genetic disorders and other traits. The nucleotide sequences are also useful as hybridisation probes, as oligomers for PCR, for chromosome and gene mapping, in the recombinant production of protein, or in generating antisense DNA or RNA. The purified polypeptide is useful for generating antibodies specific for it. The present sequence is one of the 38043 isolated cDNA/EST sequences. Note: The sequence data for this parent did not form part of the printed specification, but was obtained in electronic format directly from USPTO at
                                       used to determine the phenotype of a cell. The primers are useful for detecting a mutation in a test nucleotide sequence and also for detecting cancer, preferably colon cancer. Antibodies against the protein encoded by the nucleotide sequence can also be used in a method to detect colon cancer. The diagnostic method is non-invasive and accurate for diagnosing
                                                                                                                                                                This sequence represents a human nucleotide sequence which is differentially expressed in colon cancer cells compared to the expression levels in normal cells. The nucleotide sequence can be used as a source of primers and probes. The nucleotide sequence is useful for determining the phenotype of a cell by detecting the differential expression of the sequence relative to a normal cell. The probes derived from the sequence can also be used to determine the phenotype of cells in a sample. Probes and antibodies which hybridise to the nucleotide sequence can also be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Endege
Catino
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                31-AUG-1998;
27-JAN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO200012702-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2000-256641/22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            30-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (FARB ) BAYER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          258
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ₫.8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         l Similarity
16; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       mutation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        476
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                         eic acids and treating and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Steinmann
Dwivedi P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
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                                                                                                                                                                                                                                                                                                                                                                                             206; 345pp;
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99US-0117393P
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Ford
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94.1%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          91
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                                                                                                                                                                                                                                                                                                                                                                                                                                               for identifying therapeutic agents g cancer, especially colon cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               JH, Burgess
Lewis ME, 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        <u>ი</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 No. 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        135 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 8;
.2e+02;
les 1;
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Best Local S
Matches 16
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Best Local :
                                                                                                                                                                                                                                                                                                                                               Matches
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23-JUN-1999;
02-SEP-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention provides the coding sequences and some protein sequences of proteins associated with breast cancer in humans. These sequences can be used in the diagnosis and treatment of cancers, particularly breast tumours
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human breast cancer associated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAF17596;
  Human breast
                                                18-DEC-2001
                                                                                                                                      AAS47026 standard; cDNA; 647
                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 647 BP; 216 A; 112 C; 129 G; 189 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               tumor protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     An isolated polypeptide useful for the treatment and diagnosis of tumors e.g. breast cancer comprises at least an immunogenic portion of a breast
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2001-122627/13.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Yugiu J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             12-OCT-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Page 110; 238pp; English
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    cancer
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                                              (first entry)
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99US-00339338.
99US-00389681.
99US-00433826.
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    cDNA clone
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94.1%;
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94.1%;
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Pred. No. 5.3e+02;
0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                     Score 15.4;
Pred. No. 5.
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                                                                                                                                                                                                                                                                                                                                                                  5.3e+02
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                                                                                                                                                                                                                                                                                                                                                                                           Length 647;
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                                                                                                                                                                                                           The invention relates to isolated breast tumour proteins and nucleic concludes that encode them, including immunogenic fragments of the proteins. Also included are expression vectors expressing the proteins, transformed concells and antibodies raised against the proteins or an antigen presenting coll expressing the protein. The proteins and nucleic acids may be used in the prevention, diagnosis and treatment of diseases associated with concept cancer e.g by gene therapy. The nucleic acids and their complements may also be used as DNA probes in diagnostic assays to detect and quantitate the presence of similar nucleic acids in samples, and therefore which patients may be in need of restorative therapy. The proteins, nucleic acids and antibodies may be used in assays to identify condulators (e.g. antagonists) of breast tumour protein expression and crivity. The antibodies and antagonists may also be used as crivity. The antibodies and antagonists may also be used as crivity. The antibodies and antagonists may also be used as conducted agents for detecting the presence of the proteins in samples (e.g. by enzyme linked immunosorbant assay (ELISA)) and in other immunocuparity to the presence of the proteins acond account to the proteins of the proteins and account and activity and the presence of the proteins and account 
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SUMMARIES

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Vogelstein, B., Baker, S.J., Fearon, E.R. and Nigro, J.M.
Detection of loss of the wild-type P53 gene and kits therefor
Patent: US 5527676-A 15 18-JUN-1996;
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AR060418/c
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HUMP5302/c
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                                                                                                                                                                                                                                                                                                                18;
                                                                            1 (bases 1 to 133)
Boeke, J.D. and Brachmann, R.K.
Nucleic acids encoding p53 mutations which
                                                                                                                                                                             Sequence 22
AR060418
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata;
Mammalia; Eutheria; Primates; Catarrhini
1 (bases 1 to 102)
Lamb, P. and Crawford, L.
Characterization of the human p53 gene
Characterization of the human p53 gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  antigen; phosphoprotein;
2 of 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens phosphoprotein M13112
                                                         Patent:
                                                                      mutations
                                                                                                                    Unclassified.
                                                                                                                                 Unknown
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                                                                                                                                                                                                    AR060418
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                                                                                                                                                                                                                                                                                                             100.0%; Score 18; DB 9; llarity 100.0%; Pred. No. 7.8e+02; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                           US 5840579-A 22 24-NOV-1998;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1. .102
/organism="unknown"
/mol_type="unassigned DNA"
        /organism="unknown"
/mol_type="unassigned DNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                 /organism="Homo sapiens"
/mol_type="genomic DNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                /gene="TP53"
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                                                                                                                                                                                                                                                                                                                                                                                   note="G00-120-445"
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                                                                                                                                                                                        133 bp
US 5840579.
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122232/c
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AR128976/c
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HUMP53A02/c
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1 (base 1 to 133)

1 (base 1 to 133)

1 (base 1 to 133)

Vogelstein, B., Baker, S.J., Fearon, E.R. Detection of loss of the wild-type P33

Patent: US 5527676-A 2 18-JUN-1996;

Patent: US 5527676-A 2 18-JUN-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AR128976
Sequence
AR128976
        Human phosphoprotein
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122232
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Location/Qualifiers
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                                                                                                                                                                                                                                                                                       Unknown
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Boeke, J.D. and Brachmann, R.K.
Method for identifying suppressor mutations
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                    HUMP53A02
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22 from patent US 6183964.
                                                                                                                                                                              /organism="unknown"
/mol_type="unassigned
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/mol_type="unassigned
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100.0%; Pred. No. 7.:
tive 0; Mismatches
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100.0%; Pred. No. 7.3e+02;
tive 0; Mismatches 0;
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Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, I
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, I
Mammalia, Eutheria, Primates, Catarrhini, Hominidae,
1 (bases 1 to 140)
Preddie, E. R. and Bergmann, J. E.
AGENTS FOR THE PRE-SYMPTOMATIC DETECTION, PREVENTION
OF BREAST CANCER IN HUMANS
PATENT: WO 9807851-A 3 26-FEB-1998;
PREDDIE ENRIQUE R (CA)
                                                                                                                                                                                                                                                                                                                                                                                        140 bp
Sequence 3 from Patent WO9807851.
A69972
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1 (bases 1 to 133)
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M22882.1 GI:189465
phosphoprotein p53.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               About 10400 bp after segment 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Original source text: Homo sapiens placenta DNA Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Buchman, V.L., Chumakov, P.M., Ninkina, N.N., Georgiev, G.P.
                                                                                                                                                                                                                                                                                                                                                                             A69972.1 GI:4774443
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  h 100.0%; Score 18; DB 9; Similarity 100.0%; Pred. No. 7.3e+02;
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                                                       100.0%; Score 18; DB 6; larity 100.0%; Pred. No. 7.2e+02; Conservative 0; Mismatches 0;
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70 (2), 245-252 (1988)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /tissue_type="placenta"
order(M22881.1:397. .>412,<1. .15)
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/gene="p53"
                                                                                                                             /organism="Homo sapiens"
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/db_xref="taxon:9606"
/clone="BC532L"
                                                                                                                                                                                               Location/Qualifiers
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/number=2
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db_xref="taxon:9606"
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                                                                                                                                                      . Similarity
18; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2 (bases I to 167)
Thompson-Hehir, J., Davies, M.P.A., Green, J.A., Halliwell, N.,
Joyce, K.A., Salisbury, J., Sibson, D.R., Vergote, I. and Walker, C.
Direct Submission
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1 (bases 1 to 167)
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                                                                                                                                                  100.0%; llarity 100.0%; Conservative 0
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tissues"
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/note="in GenBank Accession Number U94788"
/replace="cc"
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/product="tumor suppressor
/protein_id="AAF36354.1"
/db_xref="GI:7107285"
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
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'gene="TP53"
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Submitted (30-NOV-1999) J. K. Douglas Cancer Research Laboratories,
Clatterbridge Cancer Research Trust, Clatterbridge Hospital,
Clatterbridge Cancer Research Trust, Clatterbridge Hospital,
Bebington, Wirral CH63 4JY, United Kingdom
Location/Qualifiers
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1 (bases 1 to 167)

Thompson-Hehir, J., Davies, M.P., Green, J.A., Halliwell, N., Joyce, K.A., Salisbury, J., Sibson, D.R., Vergote, I. and Walker, C. Novel polymerase chary, J., Sibson, D.R., Off full-coding p53 mutation detection in microdissected archival tumors Diagn. Mol. Pathol. 9 (2), 110-119 (2000)
                                                  AF209130 167 bp
Homo sapiens tumor suppressor p53
AF209130
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AF209129
                                     AF209130.1 GI:7107288
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Homo sapiens (human)
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/gene="TP53"
/note="in_GenBank Accession Number U94788"
                                                                                                                                                                                                                                                                                                                                               replace="cc"
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/codon_start=1
/product="tumor suppressor
/protein_id="AAF36355.1"
/db_xref="GI:7107287"
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mol_type="genomic DNA"
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map="17p13.1"
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Pred. No. 6.8e+02;
; Mismatches 0;
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                                                                      DNA linear PRI 27-
(TP53) gene, partial cds.
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1 (bases 1 to 167)

Thompson-Hehir, J., Davies, M.P., Green, J.A., Halliwell, N., Joyce, K.A., Salisbury, J., Sibson, D.R., Vergote, I. and Walker, C. Novel polymerase chain reaction approach for full-coding p53 mutation detection in microdissected archival tumors
Diagn. Mol. Pathol. 9 (2), 110-119 (2000)
                                                                                                                                                                                                                                                                                                                                                                         18;
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Clatterbridge Cancer Research Trust, Clatterbridge Hospital,
Bebington, Wirral CH63 47Y. United Kingdom
Location/Qualifiers
                                                                                                                                                Homo sapiens tumor suppressor
AF209131
AF209131.1 GI:7107290
Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 167)
Thompson-Hehir,J., Davies,M.P., Green,J.A., Halliwell,N., Joyce,K.A., Salisbury,J., Sibson,D.R., Vergote,I. and Walker,C.
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                                                                                                               sapiens (human)
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/note="in GenBank Accession Number U94788"
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|24. .>167
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/db_xref="GI:7107289"
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/db_xref="taxon:9606"
/chromosome="17"
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|mol_type="genomic DNA"
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essor p53
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 Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 167)

Thompson-Hahir, J., Davies, M.P., Green, J.A., Halliwell, N., Joyce, K.A., Salisbury, J., Sibson, D.R., Vergote, I. and Walker, C. Novel polymerase chain reaction approach for full-coding p53 mutation detection in microdissected archival tumors Diagn. Mol. Pathol. 9 (2), 110-119 (2000)
                                                                                                                                                                Homo sapiens (human)
Homo sapiens
                                                                                                                                                                                                                                                         167 bp DNA linear Homo sapiens cell-line Molt4 tumor suppressor p53 partial cds.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Submitted (30-NOV-1999) J. K. Douglas Cancer Research Laboratories, Clatterbridge Cancer Research Trust, Clatterbridge Hospital, Bebington, Wirral CH63 4JY, United Kingdom
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2 (bases 1 to 167)
Thompson-Hehir, J., Davies,
Joyce, K.A., Salisbury, J.,
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                                                                                                                                                                                                                          AF209132.1 GI:7107292
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<1. .>167
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /gene="TP53"
/note="in GenBank Accession Number U94788"
/replace="cc"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /codon_start=1
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/protein_id="GI:7107291"
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tissue_type="head and neck carcinomas and associated
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
/chromosome="17"
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Best Local
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                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 167)

Thompson-Hehir,J., Davies,M.P., Green,J.A., Halliwell,N., Joyce,K.A., Salisbury,J., Sibson,D.R., Vergote,I. and Walker,C. Movel polymerase chain reaction approach for full-coding p53 mutation detection in microdissected archival tumors
Diagn. Mol. Pathol. 9 (2), 110-119 (2000)
                                                                                                                                                                                                                                                                        167 bp DNA linear PRI 27-MAR-2003
Homo sapiens cell-line A431 tumor suppressor p53 (TP53) gene,
partial cds.
AF209133
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Thompson.Hehir, J., Davies, M.P.A., Green, J.A., Halliwell, N.,
Joyce, K.A., Salisbury, J., Sibson, D.R., Vergote, I. and Walker, C.
Direct Submission
2 (bases 1 to 167)
Thompson-Hehir, J., Davies, M.P.A., Green, J.A., Halliwell, N.,
Joyce, K.A., Salisbury, J., Sibson, D.R., Vergote, I. and Walker, C.
Direct Submission
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1. .167
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
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Pred. No. 6.8
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6.8e+02;
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Direct Submission
Submitted (30-NOV-1999) J. K. Douglas Cancer Research Laboratories,
Submitted (1998) S. K. Douglas Cancer Research Laboratories,
Clatterbridge Cancer Research Trust, Clatterbridge Hospital,
Bebington, Wirral CH63 4JY, United Kingdom
Location/Qualifiers
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                                                                                                                                       2 (bases 1 to 167)
Thompson-Hehir, J., Davies, M.P.A., Green, J.A., Halliwell, N.,
Joyce, K.A., Salisbury, J., Sibson, D.R., Vergote, I. and Walker, C.
                                                                                                                                                                                                                                                 Thompson-Hehir, J., Davies, M.P., Green, J.A., Halliwell, N., doyce, K.A., Salisbury, J., Sibson, D.R., Vergote, I. and Walker, C. Novel polymerase chain reaction approach for full-coding p53 mutation detection in microdissected archival tumors Diagn. Mol. Pathol. 9 (2), 110-119 (2000)
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1 (bases 1 to 167)
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/note="in GenBank Accession Number U94788"
/replace="cc"
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/map="17p13.1"
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
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Query Match
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Submitted (30-NOV-1999) J. K. Douglas Cancer Research Laboratories, Clatterbridge Cancer Research Trust, Clatterbridge Hospital, Bebington, Wirral CH63 4JY, United Kingdom
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             167 bp DNA linear PRI 27-M2
Homo sapiens cell-line HeLa tumor suppressor p53 (TP53) gene,
partial cds.
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                                                                                                                                                                                                                                                                                                  Thompson-Hehir, J., Davies, M.P., Green, J.A., Halliwell, N., Joyce, K.A., Salisbury, J., Sibson, D.R., Vergote, I. and Walker, C. Novel polymerase chain reaction approach for full-coding p53 mutation detection in microdissected archival tumors Diagn. Mol. Pathol. 9 (2), 110-119 (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 167)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /mol_type="genomic DNA;
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
/chromosome="17"
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|24. .>167
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /codon_start=1
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Submitted (30-NOV-1999) J. K. Douglas Cancer Research Laboratories, Clatterbridge Cancer Research Trust, Clatterbridge Hospital, Bebington, Wirral (H63, 477, United Kingdom
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               167 bp DNA linear PRI 27-MAR-2003
Homo sapiens cell-line HN5 tumor suppressor p53 (TP53) gene,
                                                                                                                                                                                                                                                                                                                                                                                                                   1 (bases 1 to 167)
Thompson-Hehit, J., Davies, M.P., Green, J.A., Halliwell, N.,
Thompson-Hehit, J., Davies, M.P., Green, J.A., Halliwell, N.,
Joyce, K.A., Salisbury, J., Sibson, D.R., Vergote, I. and Walker, C.
Novel polymerase chain reaction approach for full-coding p53
mutation detection in microdissected archival tumors
Diagn. Mol. Pathol. 9 (2), 110-119 (2000)
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AF209136.1
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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<1. .21
                                                         /map="17p13.1"
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/replace="cc"
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<1. .>167
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'product="tumor suppressor p53
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/gene="TP53"
                                                                                                               chromosome="17"
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RESULT 17 AF209135/c LOCUS

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Matches

KEYWORDS SOURCE

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ACCESSION DEFINITION

VERSION

REFERENCE

AUTHORS

PUBMED REFERENCE

AUTHORS

JOURNAL MEDLINE TITLE

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FEATURES

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AF210308.1 GI:7542419
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                                                                                                                                                                                                                                                                                     Submitted (01-DEC-1999) J.K. Douglas Cancer Research Laboratories,
Clatterbridge Cancer Research Trist, Clatterbridge Hospital,
Bebington, Wirral CH63 4JY, UK
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                2 ((bases 1 to 167)
Thompson-Hehir, J., Davies, M.P.A., Green, J.A., Halliwell, N.,
Joyce, K.A., Salisbury, J., Sibson, D.R., Vergote, I. and Walker,
Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Thompson-Hehir, J., Davies, M.P.A., Green, J.A., Halliwell, N., Joyce, K.A., Salisbury, J., Sibson, D.R., Vergote, I. and Walker, C. Mutation Detection Utilizing a Novel PCR Approach for Amplification of the p53 Gene from Microdissected Tissue: Application to Archival
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 167)
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                                                                                                                                                                                                                                                                                                                                                                                                                                  Tumor Samples Unpublished
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                                                                                                                                                                                               /organism="Homo sapiens"
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'gene="TP53"
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                                                                       gene="TP53"
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                                                                                                                                                                                                                                                                                                                 255 bp
Sequence 5 from Patent WO9807851.
A69974
A69974.1 GI:4774444
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Sequence 5 from Patent W09421791.
A40037
A40037.1 GI:2296203
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   l Similarity
18; Conserv
                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 255)

Preddie, E.R. and Bergmann, J.E.
AGENTS FOR THE PRE-SYMPYCMATIC DETECTION, PREVENTION AND TREATMENT OF BREAST CANCER IN HUMANS
PATENT: WO 9807851-A 5 26-FEB-1998;

PREDDIE ENRIQUE R (CA)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
1 (bases 1 to 255)
Bergmann, J. E. and Preddie, R. E.
AGENTS FOR THE PREVENTION AND TREATMENT OF BREAST CAN
Patent: FO 9421791-A 5 29-8EP-1994;
BERGMANN JOHANNA EUGENIE (DE)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 CCCGGAAGGCAGTCTGGC 18
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Similarity 100.0%; Pred. No.
18; Conservative 0; Mismatch
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ilarity 100.0%;
Conservative (
100.0%; Score 18; DE larity 100.0%; Pred. No. 6.1 Conservative 0; Mismatches
                                                                         /organism="Homo sapiens"
/mol_type="unassigned DNA"
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/clone="BC533L"
                                                                                                                                Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /organism="Homo sapiens"
/mol_type="unassigned DNA"
/db_xref="taxon:9606"
/clone="BR538.1"
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Pred. No. 6.8e+02
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thes 0;
                6.1e+02;
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                                DB 6;
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122243/c
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192481/c
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Sequence 1
192481
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Vogelstein,B., Baker,S.J., Fearon,B.R. and Nigro,J.M.
Detection of loss of the wild-type P53 gene and kits therefor Patent: US 5527676-A 13 18-JUN-1996;
Location/Qualifiers
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                                                                           Sequence 13 from patent 122243
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George,A.L. Jr., Bhatnagar,S.K. and Nazarenko,I.
Method for analyzing a nucleotide sequence
Patent: US 5728526-A 1 17-MAR-1998;
Location/Qualifiers
           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; I
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
                                 Homo sapiens (human)
                                                                                                                                                                                                                                                                                                                                                           Unknown
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Unclassified
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                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                      /organism="unknown"
/mol_type="unassigned
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /mol_type="unassigned
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US 5527676
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                      Euteleostomi;
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RESULT 26
HUMTP53B/c
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TITLE
JOURNAL
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Best Local Similarity 100.0%;
Matches 18; Conservative 0
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S Lane, D.P., Jackson, S.P., Lakin, N.D. and Smith, G.C.M.

S Lane, D.P., Jackson, S.P., Lakin, N.D. and Smith, G.C.M.

S Lane, D.P., Jackson, S.P., Lakin, N.D. and Smith, G.C.M.

S Lane, D.P., Jackson, S.P., Lakin, N.D. and Smith, G.C.M.

B patent: JP 2001510684-A 2 07-AUG-2001;

KUDOS PHARMACEUTICALS LTD

EN UP 2001510684-A/2

PD J 2001510684-A/2

PD 07-AUG-2001

PD 07-AUG-2001

PD 07-AUG-2001

PD 07-AUG-2001

PD 07-AUG-2001

PD 16-UUL-1998 UP 2000503426

PF 16-UUL-1997 GB 9714971.0

PD 16-UUL-1997 GB 9714971.0
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Human p53 cellular ti
M14695
M14695.1 GI:339815
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Patent: WO 9904266-A 4 28-JAN-1999;
SMITH GRAEME CAMERON MURRAY (GB); CANCER RES
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Similarity 100.0%;
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Key Location/Qualifiers
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/mol_type="unassigned DNA"
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
                                                cellular tumor
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                                                                PRI 14-JAN-1995
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Matches 18
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                                                                                                                                                                                                                  Unclassified.

Unclassified.

( bases 1 to 1307)

( bases 1 to 1307)

Vogelstein,B., Baker,S.J., Fearon,E.R. and Vogelstein, B., Baker, B.J., Fearon,E.R. and Vogelstein, B., Baker, B.J., Fearon,E.R. and Vogelstein, B., Baker, B.J., Fearon,E.R. and Vogelstein, B. and Vogelstein, Fearon, Fearon,
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Harris,N., Brill,E., Shohat,O.,
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I22242
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Original source text: Human transformed cell line SV-80, cDNA
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Mammalia; Eutheria; Primates;
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                                                                                  Conservative
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                                                                                                                                                                                                            /organism="unknown"
/mol_type="unassigned DNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GI:1602596
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/note="p53 antigen"
/codon_start=1
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/mol_type="mRNA"
/db_xref="taxon:9606"
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                                                                               100.0%; Score 18; DB 6; 1
100.0%; Pred. No. 3.8e+02;
ive 0; Mismatches 0;
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Pred. No. 3.9e+02;
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US 5527676.
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RESULT 29
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M14694.1
METHOD AND MEANS FOR DISRUPTION OF PAtent: WO 9741433-A 6 06-NOV-1997; CANCER RES CAMPAIGN TECH (GB)
Location/Qualifiers
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Harris,N., Brill,E., Shohat,O., Prokocimer,M., Wolf,D., Arai,N. and
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/db_xref="G1:339814"
/db:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:ychtofid=ydb:
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'db_xref="taxon:9606"
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100.0%; Pred. No. 3.8e+02;
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c antigen mRNA, complete cds.
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AR123180/c
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1 (bases 1 to 1317)

1 (bases 1 to 1317)

Halazonetis, T.D.

Modified p53 constructs which enhance DNA binding Patent: US 5847083-A 1 08-DEC-1998;

Location/Qualifiers
1. .1317
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Unclassified.

1 (bases 1 to 1317)

Halazonetis, T. and Hartwig, W.
Peptides and peptidomimetics with structural similarity
p53 that activate p53 function
Patent: US 6169073-A 1 02-JAN-2001;
                                                                                                                                                         AR123180 1317 bp
Sequence 1 from patent US 6169073.
AR123180
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AR064393
                                                                                               Unknown
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/db_xref="G1.4538507"
/db_xref="G1.4538507"
/db_xref="RMTERMEL:Cab39356"
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/translation="MEEPQSDESVEPPLSQETFSDLWKLLPENNVLSPLPSGVPSQKT
/translation="MEEPQSDESVEPPLSQETFSDLWKLLPENNVDSTPPPGTRVRAM
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PYEPPEVGSDCTTIHYNYMCNSSCMGGMNRRPILTIITLEDSSGNLLGRNSFEVRVCA
CPGRDRRTEBENLRKKGSPHHBLPPGSTTRALPNTSSSPQPKKKPLDGEYFTLQIRG
REKFEMFRELNEALELKDAQAGKEPGGSRAHSSHLKSKKGQSTSRHKKLMFKTEGPDS
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/mol_type="unassigned DNA"
/db_xref="taxon:32644"
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/mol_type="unassigned
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100.0%; Pred. No. 3.8e+02;
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.0%; Pred. No. 3.8e+02;
Mismatches 0;
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AR157699/c
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    127
                                                                                                                                                                                                                   Unclassified.

1 (bases 1 to 1317)

1 (bases 1 to 1317)

Halazonetis, T.D.

P53 proteins with altered tetramerization domains

Patent: US 5573925-A 1 12-NOV-1996;

Patent: US 5573925-A 1 10-NOV-1996;

1. 1317

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Sequence 1 f
AR157699
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Halazonetis, T. and Hartwig, W.
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                                                                                     100.0%; Score 18; DB 6; ilarity 100.0%; Pred. No. 3.8e+02; Conservative 0; Mismatches 0;
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/mol_type="unassigned
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/mol_type="unassigned
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/mol_type="unassigned
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from patent US 557
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patent US 6245886
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I89361/c
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AR218593/c
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Halazonetis,T.D.
p53 prote:
                                                                         AR218593
Sequence 1 :
AR218593
AR218593.1
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                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
AR181303
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Unclassified.
1 (bases 1 to 1317)
Halazonetis, T. and Hartwig, W.
Peptides and peptidomimetics with
                                                                                                                                                                                                                                                                                        1 (bases 1 to 1317)
Kigawa, K., Yamanaka, M., Kusumi, K., Mukai, E. and Obata, K.
Methods for targeting, enriching, detecting and/or isola
nucleic acid sequence using Reca-like recombinase
patent: US 6335164-A 8 01-JAN-2002;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   18;
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                                                                                                                                                                                                                                                          /mol_type="unassigned DNA"
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/mol_type="unassigned
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100.0%; Pred. No. 3.8e+02;
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AX057140/c
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                        Homo sapiens (human)
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Sequence 96 from Patent
AX203243
AX203243.1 GI:15392615
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 16 from Patent
AX057140
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Eukaryota; Metazoa; Chordata;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Zhang,H., Tsvetkov,L.M. and Kondo,T. Modulation of protein levels using the Patent: WO 0075184-A 16 14-DEC-2000; YALE UNIVERSITY (US)
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/db_xref="G1:1230962"
/db_xref="G1:1230962"
/db_xref="G1:1230962"
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/translation="MEEPQSDLWKLPSQETFSDLWKLPSQAMDDLM
/translation="MEEPQSDLWKLPSQETFSDLWKLPSQAMDDLM
/translation="MEEPQSDLWKLPSQAMDDLM
/translation="Meepqueppls"

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CPGRDRRTEEENLRKKGEPHHELPPGSTKRALPNNTSSSPQPKKKPLDGEYFTLQIRG
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/mol_type="unassigned DN:
/db_xref="taxon:9606"
|36. .1317
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   Craniata; Vertebrata;
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   Euteleostomi;
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Human mRNA for p53
X02469 M60950
X02469.1 GI:35209
                                                               HSP53
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PC
CC
CC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BD064183 1317 bp DNA linear PAT 27-1
Mehods for targeting, enriching, detecting and/or isolating
nucleic acid sequence using RecA-like recombinase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Rapid determination of gene structure
Patent: WO 0153529-A 96 26-UUL-2001;
Genome Therapeutics Corporation (US)
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo
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29-AUG-1996 JP 8/229061,26-DEC-1996 JP 8/3470
KIGAWA,MIKAYO YAMANAKA,KAYO KUSUMI,ERI MUKAI,KAZUAKI
                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%; Score 18; larity 100.0%; Pred. No. Conservative 0; Mismatch
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Strandedness: Dou
Topology: Linear;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        C12Q1/68
                                                                                                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
1. .1317
                                                                                                                                                                                                                                                                                                                                                            /organism="Conus quercinus"
/mol_type="genomic DNA"
/db_xref="taxon:101313"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /organism="Homo sapiens"
/mol_type="unassigned DNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                             100.0%; Score 18; DB 6; 100.0%; Pred. No. 3.8e+02;
                                      1317 bp mRNA
cellular tumor antigen.
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AZUAKI PI
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ORGANISM
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ACCESSION
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KEYWORDS
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E13558/c
LOCUS
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                                                                                                                                                                                                                                                                                                            E13558
E13558.1 GT:3252363
JF 1997206070-A/1.
Homo sapiens (human)
Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           18;
                                                                                                                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrat; Mammalia; Eutheria; Primates; Catarrhini; Hominid 1 (bases 1 to 1760)
Kawaguchi, R. and Hikichi, K. CELL-LINE ORIGINATED FROM HUMAN BONE MARROW CELLS Patent: JP 1997206070-A 1 12-AUG-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; E
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
1 (Dases 1 to 1317)
Zakut-Houri,R., Bienz-Tadmor,B., Givol,D. and Oren,M.
Human p53 cellular tumor antigen: cDNA sequence and e
                                                                                                                                                                                                                                                                                                                                                                                                                           Mutated p53 gene.
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                                      12-AUG-1997
31-JAN-1996 JP 1996037400
KAWAGUCHI RYUJI, HIKICHI KAZUMASA
C12N5/06,C07H21/04,C12N9/08//C12N15/09,C12Q1/68,(C12N5/06,
                                                                                                                     Homo sapiens (human)
JP 1997206070-A/1
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ilarity 100.0%;
Conservative (
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(C12N9/08, C12R1:91);
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384. .398
/note="direct repeat 1"
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/db_xref="G1:35210"
/db_xref="G2:35210"
/db_xref="G3:35210"
/db_xref="G4A:264637"
/db_xref="SWISS-PROT:P04637"
/db_xref="G4A2630"
/db_xref="SWISS-PROT:P04637"
/db_xr
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/mol_type="mRNA"
/db_xref="taxon:9606"
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1. .1317
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Pred. No. 3.8e+02;
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                                                                                                                                                                                                                                                                          Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
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ORGANISM
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E13737/c
                           ORIGIN
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VERSION
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Best Local
                                                                                                              source
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                                                                                                                                              AUTOANTIBODY DETECTING METHOD FOR P53 PROTEIN
PATENTIBODY DETECTING METHOD FOR P53 PROTEIN
POS HOMO SAPELERS (CRP
POS HOMO SAPELERS (P1996124123
PI SUBANO KOKICHI, SAOTOME KEIICHI
PC GOIN3/53,COTK14/47,COTK16/42,GO1N33/564;
PT GOIN3/53,COTK14/47,COTK16/42,GO1N33/564;
PT GOUNGES: Double;
PT GOUNGES: Double;
PT Key
PT SOURCE
PT 1. .1760
PT CDS
PT 1. .214
PT 1. .215
PT 1. .216
PT 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 1760) Sugano, K. and Saotome, K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              E13737 1760 bp DNA 1760 bp DNA cDNA encoding human p53 protein which is as
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llarity 100.0%;
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                                              /organism="Homo sapiens"
/mol_type="genomic DNA"
/db_xref="taxon:9606"
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tion replace(957,'g')
tion replace(1317,'a').
Location/Qualifiers
                                                                                                                                 location/Qualifiers
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Linear;
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/cell type='bone marrow-derived
_line='CMC-1'
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Pred. No. 3.5e+02;
; Mismatches 0;
                                                                                                                                                                                                                    p53 protein which
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                                                                                                                                                                                                                                        RESULT 44
HUMP53T/c
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antigen; antigen p53.
Homo sapiens (human)
Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Nammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 1760)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 HENRI BURE
PC C12Q1/
A01K67/027,
PC A01K48
CC Strand
CC Topolo
CC p53 ce
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Diagnosis method and reagents
patent: JP 200152241-A 13 13-NOV-2001;
ROYAL NETHERLANDS ACADEMY OF ARTS AND SCIENCES, ERASMUS UNIVERSITY
ROYTERDAM, UNIVERSITY OF UTRECHT
S Homo sapiens (human)
PN JP 2001522241-A/13
PD 13-NOV-2001
PF 02-APR-1998 UP 1998542545
PR 10-APR-1997 US 60/043163
PI FREDERIK W VAN LEEUWEN, FRANKLIN G GROSVELD, JOHANNES PETER PI
HENDI HITMAGE
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                                                                                                                     Human p53 cellular tumor antigén mRNA, complete cds K03199
K03199.1 GI:189478
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens (human)
                                                                                                                                                                            HUMP53T
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1 (Dases 1 to 1760)

Leeuwen, F.W.V., Grosveld, F.G. and Burbach, J.F.H.
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BD084820
Harlow,E
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                                                                                                                                                                                                                                                                                          CCCGGAAGGCAGTCTGGC 18
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llarity 100.0%; Pred. No. 3.
Conservative 0; Mismatches
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Topology: Linear;
p53 cellular tumor antigen gene; Genbank accession
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                                                                                                                                                                                                                                                                                                                                                                                                                    /organism="Homo sapiens"
/mol_type="genomic DNA"
/db_xref="taxon:9606"
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Pred. No. 3.5e+02;
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3.5e+02;
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A draft entry and sequence for [2],[1] were kindly submitted via electronic mail by E.Harlow 17-OCT-1985. [1] reports that positions 1-79 may be a cloning artifact generated by ligating a second, unrelated CDNA to the 5' end of the p53 cDNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Harlow,E., Williamson,N.M., Ralston,R., Helfman,D.M. and Adams,T.E. Molecular clonking and in vitro expression of a CDNA clone for human cellular tumor antigen p53 cell. Biol. 5 (7), 1601-1610 (1985)
                                                                                                                                                                                Expression vector AdSCMV-p53 tumor supperpression cassette, complete sequence. AF192534
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              McBride,O.W., Merry,D. and Givol,D.
The gene for human p53 cellular tumor antigen is loc
chromosome 17 short arm (17p13)
Proc. Natl. Acad. Sci. U.S.A. 83 (1), 130-134 (1986)
                                                          Expression vector Ad5CMV-p53
Expression vector Ad5CMV-p53
artificial sequences; vectors
1 (bases 1 to 2355)
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  Generation
                                  Zhang, W.W., Fang, X., Branch, C.D., Mazur, W., French, B.A. and
                                                                                                                                                               AF192534.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RERFEMFRELNEALELKDAQAGKEPGGSRAHSSHLKSKKGQSTSRHKKLMFKTEGPDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /protein_id="AAA59989.1"
/db_xref="GI:189479"
/db_xref="GDB:G00-120-445"
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mol_type="mRNA"

db_xref="taxon:9606"
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and identification of
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100.0%; Pred. No. 3.5e+02;
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                                                                                                                                                                                                                          tumor suppressor
  recombinant adenovirus
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AB082923/c
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Homo sapiens mRNA for P53, cc
AB082923
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Submitted (06-OCT-1999) Oncology, Rh. Bay Center Place, Hayward, CA 94545,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     High-efficiency gene transfer and high-level expression of wild-type p53 in human lung cancer cells mediated by recom
                                                                                                                                                     Homo sapiens (human)
Homo sapiens
Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          liposome-mediated transfection and I BioTechniques 15 (5), 868-872 (1993)
                                Mutated p53 gene encodes a nonmutated epitope HLA-B*4601-restricted and tumor cell-reactive Cancer Res. 63 (4), 854-858 (2003)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Direct Submission
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               94092464
                                                                                     Azuma, K., Shichijo, S
Koike, K. and Itoh, K.
                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
                    22479085
                                                                                                                                                                                                                          AB082923.1
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ang, W.W., Fang, X., Mazur, W.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (bases 1 to 2355)
                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
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RERFEMFRELNEALELKDAQAGKEPGGSRAHSSHLKSKKGQSTSRHKKLMFKTEGPDS
D"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /translation="MEEPQSDPSVEPPLSQETFSDLWKLLPENNVLSPLPSQAMDDLM LSPDDIEQWFTEDFGFDEAPRAPAPAPAPAPAPAPAPAPAPAPSSVPSQKT YQGSYGFRLGFLHSGTAKSVTCTYSPALNKMFCQLAKTCPVQLWVDSTPPFGTRVRAM AIYKQSQHMTEVVRCCHHERCGDSDGLAPPQHLIRVEGNLRVEYLDDRNFFRLSVVV PYEPPEVGSDCTTIHYNYMCNSSCMGGMNRRPILTIITLEDSSGNLLGRNSFEVRVCA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /noTe="p53 expression cassette of RPR/INGN 201 readenovirus vector. Artificial sequence; contains promoter and the SV40 polyadenylation signal"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /gene="p53"
921. .2102
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /gene="p53"
/codon_start=1
/product="tumor_suppressor_protein_p53"
/protein_id="AAF64408.1"
/db_xref="GI:7595312"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /organism="Expression vector AdSCMV-p53"
/mol_type="genomic DNA"
/db_xref="taxon:120786"
                                                                                                    Shichijo, S., Maeda, Y., Nakatsura, T.,
                                                                                                                                                                                                                          GI:23491728
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                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 18; DB 12;
Pred. No. 3.3e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     and Roth, J.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Rhone-Poulenc
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                                                                                                                                                                                                                                                                             linear
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                                                    recognized by CTLs at tumor
                                                                                                       Nonaka, Y., Fujii, T.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Rorer Gencell,
                                                                                                                                                          Euteleostomi;
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REFERENCE

(bases 1 to 2451)

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RESULT 47
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AUTHORS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM
                   TITLE
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Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi; Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi; Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo.

CE 1 (bases 1 to 2508)

CE 1 (bases 1 to 2508)

Strausberg, R. L., Feingold, E. A., Grouse, L. H., Derge, J.G., R. Strausberg, R. D., Collins, F. S., Wagner, L., Shenmen, C. M., Schuler, G. D., Altechul, S. F., Zeeberg, B., Buetow, K. H., Schaefer, C. F., Bhat, N. K., Hopkins, R. F., Jordan, H., Moore, T., Max, S. I., Wang, J., Hsieh, F., Diatchenko, L., Marusina, K., Farmer, A. A., Rubin, G. M., Hong, L., Stapleton, M., Soares, M. B., Bonaldo, M. F., Casavant, T. L., Scheetz, T. E., Brownstein, M. J., Usdin, T. B., Toshiyuki, S., Scheetz, T. E., Brownstein, M. J., Usdin, T. B., Toshiyuki, S., Carninci, P., Prange, C., Raha, S. S., Loquellano, N. A., Peters, G. J., Abramson, R. D., Mullahy, S. J., Bosak, S. A., McEwan, P. J., K., Hale, S., Garcia, A. M., Gay, L. J., Hulyk, S. W., Worley, K. C., Hale, S., Garcia, A. M., Gay, L. J., Hulyk, S. W., Villalon, D. K., Muzny, D. M., Sodergren, E. J., Lu, X., Gibbs, R. A., Fahey, J., Helton, E., Ketteman, M., Madan, A., Rodrigues, S., Sanchez, A., Mhiting, M., Madan, A., Young, A. C., Shevchenko, Y., Bouffard, G. G., Blakesley, R. W., Touchman, J. W., Green, E. D., Dickson, M. C., Rodriguez, A. C., Grimwood, J., Schmutz, J., Myers, R. M., Schmert, A., Schein, J. E., Jones, S. J. and Marra, M. A., Generation and initial analysis of more than 15,000 full-length
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BC003596 2508 bp mRNA linear PRI 03-OCT-200 Homo sapiens tumor protein p53 (Li-Fraumeni syndrome), mRNA (cDNA clone MGC:646 IMAGB:3544714), complete cds.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Shichijo, S. and Itoh, K. Direct Submission
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/protein id="BAC16799.1"
/db_xref="G1:23491729"
/translation="MEBPOSDPSVEPPLSQETFSDLWKLLPENNVLSPLPSQAMDDLM
LSPDDIEQWFTEDPGPDEAPRWFEAAPRVAPAFAAPTFAAPAFAPSWFLSSSVPSQKT
YQGSVGFRLGFLHSGTAKSVTCTYSPALIKMFCQLAKTCFVQLWVDSTPPPGTRVRAM
AIYKQSQMTEVVRRCPHHERCSDSDGLAPPQHIIRVEGNURQEWYETRHSWVV
FYEPPEWGSDCTTIHYNYMCNSSCMGGMNRRFILTIITTEDSGGNLLGENFFUHVCA
CPGRDRRTEEBNLRKKGEPHHELPPGSTKRALSNNTSSSPQPKKFPLDGEYFTLQIRG
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/codon_start=1
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mol_type="mRNA"
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Pred. No. 3.2e+02;
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Matches 18
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This clone was selected for full length sequencing because it passed the following selection criteria: matched mRNA gi: 8400737.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Steven Jones, Jennifer Asano, Ian Bosdet, Yaron Butterfield, Susanna Chan, Readman Chiu, Chris Fjell, Erin Garland, Ran Guin, Letticia Hsiao, Martin Krzywinski, Reta Kutsche, Oliver Lee, Soo Sen Lee, Victor Ling, Carrie Mathewson, Candice McLeavy, Steven Ness, Pawan Pandoh, Anna-Tiisa Prabhu, Parvaneh Saeedi, Jacqueline Schein, Duane Smailus, Michael Smith, Lorraine Spence, Jeff Stott, Michael Thorne, Miranada Tsai, Natasja van den Bosch, Jill Vardy, George Yang, Scott Zuyderduyn, Marco Marra.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Email: cgapbs-r@mail.nih.gov
Tissue Procurement: DCTD/DTP
cDNA Library Preparation: Rubin Laboratory
cDNA Library Preparation: Rubin Laboratory
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Genome Sequence Centre,
BC Cancer Agency, Vancouver, BC, Canada
info@bcgsc.bc.ca
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Submitted (20-FEB-2001) National Institutes of Health, Mammalian Gene Collection (MGC), Cancer Genomics Office, National Cancer Institute, 31 Center Drive, Room 11A03, Bethesda, MD 20892-2590,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Contact: MGC help desk
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Strausberg, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          human and mouse cDNA sequences Proc. Natl. Acad. Sci. U.S.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /codon start=1
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/protein id="ak#03596,1"
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/db_xref="GI:13097807"
/db_xref="GI:157"
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                                                                                                                                                                                                                                                 /db_xref="CDD:pfam00870"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /tissue type="Kidney, renal cell adenocarcinoma"
/clone lib="NIH_MGC_14"
/lab_host="DH10B-R"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /organism="Homo sapiens"
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/cione="MGC:646 IMAGE:3544714"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            note="synonyms: p53, TRP53"
/db_xref="LocusID:7157"
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                                               100.0%;
                                               Score 18; DB 9;
Pred. No. 3.2e+02;
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AF307851/c
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DEFINITION
                                                                                                                                                                                   RESULT 49
HOMOTSP1/c
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1 (bases 1 to 2521)

Chang,N.S., Pratt,N., Heath,J., Schultz,L., Sleve,D., Carey,G.B. and Zevotek,N.

Hyaluronidase induction of a WW domain-containing oxidoreductase that enhances tumor necrosis factor cytotoxicity

J. Biol. Chem. 276 (5), 3361-3370 (2001)
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 3407)
                                         Homo sapiens (human)
Homo sapiens
                                                                                                       Homo sapiens tumor suppressor
through 9.
AF136270.1 GI:4732144
                                                                                                                                                                                                                                                                                                             18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Submitted (21-SEP-2000) Laboratory of Molecular Immunology, Guthrie Research Institute, 1 Guthrie Square, Sayre, PA 18840, USA Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AF307851
AF307851.1
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                                                                                                                                                                    HOMOTSPI
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Direct Submission
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                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                           /codon_start=1
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CPGRDRRTEEENLRKKGEPHHELPPGSTKRALPNTSSSPQPKKKPLDGEVFTLQIRG
CPGRDRRTEEENLRKKGEPHHELPPGSTKRALPNTSSSPQPKKKPLDGEVFTLQIRG
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D"
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/db_xref="dbEST:AI243172"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    organism="Homo sapiens"

mol type="mRNA"
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Pred. No. 3.2e+02;
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                                                                                                                                                    (P53) gene,
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HSM059JP1/c
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Local Similarity 100.0%;
hes 18; Conservative (
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AF135120
AF135120.1
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                                                                                                                        Homo sapiens
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                                                                                                                                       Homo sapiens (human)
                                               Mirzayans,R.
                                                                                                                                                                                                                                                HSM059JP1
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/gene="P53"
/number=2
407. .428
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/gene="P53"
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/gene="P53"
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/gene="P53"
/number=7
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/gene="P53"
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/gene="P53"
/number=5
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522. .800
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Submitted (20-MAR-1999) Biology, Basell Avenue, Upton, NY 11973-5000,
Location/Qualifiers
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                                                                                     Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Mammalia, Eutheria, Primates, Catarrhini, Hominidae, 1 (bases 1 to 3423)
Allalunis-Turner, M.J., Barron, G.M., Day, R.S. III, Dol
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Isolation of two cell lines from a human malignant glioma specimen differing in sensitivity to radiation and chemotherapeutic drugs Radiat. Res. 134 (3), 349-354 (1993)
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/db xref="kATCC:CCL-185"
/db xref="taxon:9606"
/cell_line="A549"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note="A549 cells express wild-type
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Pred. No. 2.9e+02;
; Mismatches 0;
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                                                                                             Dobler, K.D. and
                                                                                                                                                                   Euteleostomi;
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Anderson, C.W. and Allalunis-Turner, M.J.
Human TP53 from the malignant glioma-derived cell lines M059J and
M059K has a cancer-associated mutation in exon 8
Radiat. Res. 154 (4), 473-476 (2000)
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Anderson,C.W., Kieleczawa,J. and Allalunis-Turner,M.J.
Direct Submission
Submitted (16-MAR-1999) Biology, Brookhaven National Laboratory, 50
Bell Avenue, Upton, NY 11973-5000, USA
Bell Avenue, Upton, NY 11973-5000, USA
1 .3423
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Minimum DB seq length: Maximum DB seq length:
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Listing first 100 :
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   Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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Sequence 35, Appl

Sequence 22, Appl

Sequence 22, Appl

Sequence 6, Appli

Sequence 1, Appli

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Sequence 9, Appli

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Sequence 16, Appli

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Result No.

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Sequence 35, Application US/09848868
Publication No. US20030166588A1
GENERAL INFORMATION:
APPLICANT: Iversen, Patrick L.
APPLICANT: Hudziak, Robert
TITLE OF INVENTION: Method
FILLE OF INVENTION: Method
CURRENT APPLICATION NUMBER: US/09/848,868
CURRENT APPLICATION NUMBER: US/09/848,868
CURRENT FILING DATE: 2001-05-04
PRIOR APPLICATION NUMBER: US 60/202,376
PRIOR APPLICATION NUMBER: US 50/202,376
PRIOR FILING DATE: 2000-05-04
NUMBER OF SEQ ID NOS: 54
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 35
LENGTH: 18
LENGTH: 18
TYPE: DNA
ORGANISM: Artificial Sequence
                                                                     APPLICANT: Iversen, Patrick L.
APPLICANT: Hudziak, Robert
TITLE OF INVENTION: Splice-Region Antisense Composition and
TITLE OF INVENTION: Method
FILE REFERENCE: 0450-0037.30
FILE REFERENCE: 0450-0037.30
CURRENT APPLICATION NUMBER: US/09/848,868
CURRENT FILING DATE: 2001-05-04
FRIOR APPLICATION NUMBER: US/09/848,868
FRIOR APPLICATION NUMBER: US/09/848,868
FRIOR FILING DATE: 2001-05-04
PRIOR FILING DATE: 2001-05-04
PRIOR FILING DATE: 2001-05-04
NUMBER OF SEQ ID NOS: 54
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 36
LENGTH: 36
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OTHER INFORMATION: antisense
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OTHER INFORMATION: antisense
                                      ORGANISM: Artificial Sequence
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No. US20030166588A1
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Best Local Similarity
Matches 18; Conserv
                                                                                                   Sequence 6, Application US/10397635
Publication No. US20040009604A1
GENERAL INFORMATION:
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Best Local Similarity
APPLICANT: ZHANG, XIAOLIU
APPLICANT: FU, XUPING
TITLE OF INVENTION: POTENT ONCOLYTIC HERPES SIMPLEX VIRUS FOR CANCER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEX: <Unknown>
INFORMATION FOR SEQ ID NO: 22:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FASTSED for Windows Version
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/776,695
FILING DATE: 26-Feb-2001
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
OPERATOR OF THE CONTROL OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Brachmann, Rainer
TITLE OF INVENTION: NUCLEIC ACIDS ENCODING P53
MUTATIONS WHICH SUPPRESS P53 CANCER MUTA- TIONS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Kagan, Sarah A
REGISTRATION NUMBER: 32141
REFERENCE/DOCKET NUMBER: 01107.03170
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-508-9100
TELEFAX: 202-508-9299
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: 08/795,006 FILING DATE: 2001-03-28 ATTORNEY/AGENT INFORMATION:
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MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY: linear SEQ ID NO: 22:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Boeke, Jef
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 133 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SSPONDENCE ADDRESS:
ADDRESSEE: Banner & Witcoff
STREET: 1001 G Street, NW
CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: USA
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Pred. No.
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TLE OF INVENTION:

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; ORGANISM: Homo sapiens
US-10-397-635-6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
APPLICANT: Yuan, Zhi-Min
APPLICANT: Gu, JiJie
APPLICANT: Gu, JiJie
TITLE OF INVENTION: Inhibition of p53 Degradation
FILE REFERENCE: 21508-044
                                                                                                                                                                                                                                                                       Sequence 1, Application US/09829922
Publication No. US20030171537A1
GENERAL INFORMATION:
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Best Local .
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CURRENT APPLICATION NUMBER: US/10/397,635
CURRENT FILING DATE: 2003-03-26
PRIOR APPLICATION NUMBER: 60/367,788
PRIOR FILING DATE: 2002-03-27
PRIOR APPLICATION NUMBER: 60/410,024
PRIOR APPLICATION NUMBER: 60/410,024
PRIOR FILING DATE: 2002-09-11
NUMBER OF SEQ ID NOS: 14
                                                                                                                                                                                                                                                                                                                                         -09-829-922-1/c
                                                                                                                                                               APPLICANT: Halazonetis, Thanos
APPLICANT: Hartwig, Wolfgang
TITLE OF INVENTION: Peptides and peptidomimetics with
TITLE OF INVENTION: structural similarity to human p53 that activate p53
TITLE OF INVENTION: function
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/09/732,384
CURRENT FILING DATE: 2000-12-07
PRIOR APPLICATION NUMBER: 60/169,816
PRIOR FILING DATE: 1999-12-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        09-732-384-8/c
CURRENT APPLICATION NUMBER: US/09/829,922
CURRENT FILING DATE: 2001-04-11
PRIOR APPLICATION NUMBER: 08/894,327
PRIOR FILING DATE: 1997-12-04
PRIOR APPLICATION NUMBER: pctus96/01535
PRIOR FILING DATE: 1996-02-16
PRIOR APPLICATION NUMBER: 08/392,542
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQ ID NOS: 10 SOFTWARE: PatentIn Ver. :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         equence 8, Application US/09732384 atent No. US20020132977A1
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                                                                                                                                                    FILE REFERENCE: 2973.19998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: DNA
ORGANISM: Homo sapiens
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Pred. No. 7
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7.2;
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ORGANISM: Homo sapiens SEQUENCE DESCRIPTION: SEQ ID NO: US-10-160-290-1
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NUMBER OF SEQ ID NOS: 35
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 1, Application US/10160290 Publication No. US20030124557A1 GENERAL INFORMATION:
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ORGANISM: Homo sapiens
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Fatentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/160,290
FILING DATE: 04-Jun-2002
CLASSIFICATION - CUNHOWN-
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/685,027
FILING DATE: 10-Oct-2000
APPLICATION NUMBER: 08/392,542
FILING DATE: CUNHOMN-
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CORRESPONDENCE ADDRES
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Hartwig, Wolfgang
TITLE OF INVENTION: Peptides nad Peptidomimetics with
Structural Similarity to Human p53 That Activate
                                                                        ORIGINAL
                                                                                       MOLECULE TYPE: cDNA
                                                                                                                                                                                                                                                                                NAME: POSOTSKE, LAUTENCE H.
REGISTRATION NUMBER: 34,698
REFERENCE/DOCKET NUMBER: 0486.48439
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
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18; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE: Banner, Birch, M
STREET: 1001 G Street, N.W.
CITY: Washington, D.C.
                                                                                                                                  TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                             ENGTH: 1317 base pairs
                                                                                                                                                                                                                                                              TELEPHONE: 202
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                                                                        SOURCE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 McKie & Beckett
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Query Match 100.0%; Score 18; DB Best Local Similarity 100.0%; Pred. No. 7.2; Matches 18; Conservative 0; Mismatches

DB 14; Length 1317;

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; ORGANISM: Homo sapien
US-09-849-602-9
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US-09-849-602-9/c
                                                                                                                                                                                                                                SOFTWARE: PatentIn version 3.1 SEQ ID NO 4
                                                                          Matches
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Publication No. US20040005574A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
APPLICANT: Scanlan, Matthew J.
APPLICANT: Old, Lloyd J.
APPLICANT: Stockert, Elisabeth
APPLICANT: Chen, Yao-Tseng
TITLE OF INVENTION: Colon Cancer Antigen Panel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 9, Application US/09849602
Publication No. US20030165834A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Guarente, Leonard
APPLICANT: Vaziri, Homayoun
APPLICANT: Imai, Shin-Ichiro
                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: US 60/303,456
PRIOR FILING DATE: 2001-07-06
PRIOR APPLICATION NUMBER: US 10/NNN,NNN
PRIOR FILING DATE: 2002-07-05
PRIOR APPLICATION NUMBER: US 60/303,370
PRIOR FILING DATE: 2001-07-06
                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/10/191,121
CURRENT FILING DATE: 2002-07-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILE REFERENCE: L0461/7105(JRV)
CURRENT APPLICATION NUMBER: US/09/849,602
CURRENT FILING DATE: 2001-05-04
                                                                                                                                                                                                                                                                        NUMBER OF SEQ ID NOS: 18
                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: Sir2 and p53 Deacetylation FILE REFERENCE: 13407-015002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: PatentIn version 3.0
                                                                                                                                                                  LENGTH: 1760
TYPE: DNA
ORGANISM: Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ID NO 9
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                                                                                               Local
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                                 1 CCCGGAAGGCAGTCTGGC 18
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                                                                                           h 100.0%; Score 18; DB Similarity 100.0%; Pred. No. 6.9;
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                                                                          Conservative
                                                                        0,
                                                                        Mismatches
                                                                                                                DB 15;
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RESULT 10

INVENTION: TARGETING PROTEINS TO DELIVER THERAPEUTIC OR DIAGNOSTIC

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RESULT 12
US-10-430-503-16/c
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                                                                                                                                                                                                                                                                                                                                                                                                                     ; TYPE: DNA; Homo sapiens; ORGANISM: Homo sapiens US-10-429-802-25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 11
US-10-429-802-25/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-10-062-674-2065/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: ZOU, YIYU

APPLICANT: ZOU, YIYU

TITLE OF INVENTION: BIPARTITE T-CELL FACTOR (TCF)-RESPONSIVE PROMOTER FILE REFERENCE: UTSC.'752US

CURRENT APPLICATION NUMBER: US/10/429,802

CURRENT FILING DATE: 2003-05-05

PRIOR APPLICATION NUMBER: 60/377,672

PRIOR FILING DATE: 2002-05-03

NUMBER: OF SEQ ID NOS: 54

SOFTWARE: PATENTIN VEY. 2.1

SEQ ID NO 25

LENGTH: 2521

TYPE: NUMBER: 2012
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: PERL Program
SEQ ID NO 2065
LENGTH: 2209
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Publication No. US20040005559A1
GENERAL INFORMATION:
                                                                                                        GENERAL INFORMATION:
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Best Local (
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                                                APPLICANT: HUNG, MIEN-L...
**POTTCANT: LAN, KENG-LI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: US 09/625,102
PRIOR FILING DATE: 2000-07-24
NUMBER OF SEQ ID NOS: 2217
                                            APPLICANT:
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CURRENT FILING DATE: 2002-01-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Loring, Jeanne F.; Kaser, Matthew R.
TITLE OF INVENTION: MARKERS OF NEURONAL DIFFERENTIATION AND MORPHOGENESIS
FILE REFERENCE: PA-0026-1 CIP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Homo sapiens
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                                                                                                                                                                                                                                                                                                                            Local Similarity 100.0%; Score 18; local Similarity 100.0%; Pred. No. les 18; Conservative
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OU-YANG, FU
LIU, JAW-CHING
LAN, KENG-HSIN
                                                                                                                          Application US/10430503
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APPLICANT: Brachmann, Rainer
TITLE OF INVENTION: ENGINEERED OPEN READING FRI
FILE REFERENCE: 004255.00008
CURRENT APPLICATION NUMBER: US/10/077,176
CURRENT FILING DATE: 2002-02-19
NUMBER OF SEQ ID NOS: 71
SOFTWARE: FRESEQ for Windows Version 4.0
SEQ ID NO 58
LENGTH: 2629
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US-10-077-176-58/c
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FILE REFERENCE: UTSC:797US
CURRENT APPLICATION NUMBER: US/10/430,503
CURRENT FILING DATE: 2003-05-06
PRIOR APPLICATION NUMBER: 60/383,063
PRIOR FILING DATE: 2002-05-06
NUMBER OF SEQ ID NOS: 42
SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                             Sequence 58, Application US/10077176
Publication No. US20030175862A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: PatentIn Ver. 2.1 SEQ ID NO 1
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LENGTH: 2521
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Publication No. US20040028654A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: PROTAMINE-ADENOVIRAL VECTOR COMPLEXES TITLE OF INVENTION: AND METHODS OF USE FILE REFERENCE: INRP:097US CURRENT APPLICATION NUMBER: US/10/391,068 CURRENT FILING DATE: 2003-03-24 PRIOR APPLICATION NUMBER: 60/366,846 PRIOR APPLICATION NUMBER: 60/366,846 PRIOR PILING DATE: 2002-03-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQ ID NOS: 32
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: JI, LIN
APPLICANT: ROTH, JACK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: DNA ORGANISM: Homo sapiens FEATURE:
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME/KEY: CDS
LOCATION: (252)..(1433)
                    TYPE: DNA
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les 18; Conservative
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Similarity 100.0%;
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Pred. No.
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Pred. No.
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RESULT 17 US-10-077-176-61/c

Sequence 61, Application US/10077176
Publication No. US20030175862A1
GENERAL INFORMATION:

APPLICANT: Brachmann, Rainer
TITLE OF INVENTION: ENGINEERED OPEN READING FRAME FOR P53

FILE REFERENCE: 004255.00008

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FILE REFERENCE: 004255.00008

CURRENT APPLICATION NUMBER: US/10/077,176

CURRENT FILING DATE: 2002-02-19

NUMBER OF SEQ ID NOS: 71

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 60

LENGTH: 2629

TYPE: DNA

ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Brachmann, Rainer
TITLE OF INVENTION: ENGLINEERED OPEN READING FR.
FILE REFERENCE: 004255.00008
CURRENT APPLICATION NUMBER: US/10/077,176
CURRENT FILING DATE: 2002-02-19
NUMBER OF SEQ ID NOS: 71
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 59
LENGTH: 2629
 片
                                                                                                                             US-10-077-176-60
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US-10-077-176-60/c
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                                                                                                                                                                                                                                                                                Sequence 60, Application US/10077176
Publication No. US20030175862A1
GENERAL INFORMATION:
APPLICANT: Brachmann, Rainer
TITLE OF INVENTION: ENGINEERED OPEN READING FRAME FOR P53
                                                              Matches
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
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Publication No. US20030175862A1
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                                                                                           Query Match
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243 CCCGGAAGGCAGTCTGGC 226
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                                                                              Similarity
                              CCCGGAAGGCAGTCTGGC 18
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                                                              Conservative
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                                                                            100.0%; Score 18; DB : 100.0%; Pred. No. 6.6;
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100.0%; Pred. No. 6.6;
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Pred. No.
                                                              Mismatches
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CURRENT APPLICATION NUMBER: US/10/077,176

2002-02-19

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APPLICANT: Aune, Thomas M
APPLICANT: Olsen, Nancy J
TITLE OF INVENTION. Method for Predicting Autoimmune Disease
FILE REFERENCE: 1242/68
CURRENT APPLICATION NUMBER: US/10/439,388
CURRENT FILING DATE: 2003-05-16
PRIOR APPLICATION NUMBER: US 60/381,055
PRIOR APPLICATION NUMBER: US 60/381,055
PRIOR FILING DATE: 2002-05-16
NUMBER OF SEQ ID NOS: 70
SOFTMARE: Patentin version 3.2
SEQ ID NO 16
LENGTH: 3629
TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-10-439-388-16/c
; Sequence 16, Application US/10439388
; Publication No. US20030228617A1
; GENERAL INFORMATION:
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; ORGANISM: Homo sapiens
US-10-077-176-61
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; ORGANISM: Homo sapiens
US-10-439-388-16
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                                   US-10-165-216-3
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Best Local Sim
Matches 18;
                                                                                                                                                                                        Sequence 3, Application US/10165216
Publication No. US20030228675A1
GENERAL INFORMATION:
APPLICANT: Otterness, Diane M.
APPLICANT: Abraham, Robert T.
TITLE OF INVENTION: ATM Related Kinase ATX, Nucleic Acids
TITLE OF INVENTION: Encoding Same and Methods of Use
FILE REFERENCE: P-LJ 5222
CURRENT APPLICATION NUMBER: US/10/165,216
CURRENT FILING BATE: 2002-06-06
NUMBER OF SEQ ID NOS: 11
Query Match
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Best Local S
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                                                                                                                                       SEQ ID NO 3
LENGTH: 2629
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NUMBER OF SEQ ID NOS:
SOFTWARE: FastSEQ for
                                                                                                                                                                                                                                                                                                                                                                                    10-165-216-3/c
                                                                                                                                                                       NUMBER OF SEQ ID NOS:
SOFTWARE: FastSEQ for
                                            FEATURE:
NAME/KEY: CDS
LOCATION: (252)...(1433)
                                                                                                    ORGANISM: Homo sapiens
                                                                                                                       TYPE: DNA
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 Score 18;
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CURRENT APPLICATION NUMBER: US/09/918,995
CURRENT FILING DATE: 2001-07-30
PRIOR APPLICATION NUMBER: US/09/235,076
PRIOR FILING DATE: 1999-01-20
NUMBER OF SEQ ID NOS: 38054
SOFTWARE: Fast-SEQ for Windows Version 3.0
SEQ ID NO 36880
LENGTH: 436
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US-09-918-995-36880/c
Sequence 36880, Application US/09918995
; Publication No. US20030073623A1
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; ORGANISM: Homo sapiens
US-10-159-563-188
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                                                            Query Match
Best Local Similarity
Matches 17; Conserv
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SEQ ID NO 188
LENGTH: 2629
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Matches 18; Conserv
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APPLICANT: Ringner, Markus
APPLICANT: Ringner, Markus
APPLICANT: Peterson, Carsten
APPLICANT: Melter, Paul
TITLE OF INVENTION: SELECTIONS OF GENES AND METHODS OF USING THE SAME FOR
TITLE OF INVENTION: DIAGNOSIS AND FOR TARGETING THE THERAPY OF SELECT CANCERS
FILE REFERENCE: 11613.56US11
CURRENT APPLICATION NUMBER: US/10/159,563
CURRENT FILING DATE: 2002-12-09
PRIOR APPLICATION NUMBER: US 10/133,937
PRIOR FILING DATE: 2002-04-25
NUMBER OF SEQ ID NOS: 444
                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: NOVEL NUCLEIC ACID SEQUENCES OBTAINED TITLE OF INVENTION: FROM VARIOUS CDNA LIBRARIES FILE REFERENCE: 20411-756
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RESULT 22 US-10-066-543-113/c

Sequence 113, ; Publication No

Application US/10066543

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US-09-764-847-1845/c
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; ORGANISM: Homo sapiens
US-09-764-847-1845
                                                                                                                                                                                                      Prior application data removed - consult PALM or file wrapper NUMBER OF SEQ ID NOS: 2003
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 1845
LENGTH: 942
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CURRENT FILING DATE: 2002-01-31
NUMBER OF SEQ ID NOS: 3417
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 113
                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 1845, Application US/09764847 Patent No. US20020132767A1
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TITLE OF INVENTION: Nucleic Acids, Proteins,
FILE REFERENCE: PC009
                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/09/764,847
CURRENT FILING DATE: 2001-01-17
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NAME/CRY: misc_feature

LOCATION: 13, 45, 386, 545, 580, 582, 584, 590, 591, 603, 606, 609, LOCATION: 610, 616, 621, 623, 624, 625, 629, 630, 646, 652, 670, 672, LOCATION: 674, 676, 688, 700, 709, 710, 744, 753, 756, 758, 759, 763, LOCATION: 786, 796, 823, 851, 876, 885

COTHER INFORMATION: n = A,T,C or G
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TYPE: DNA
ORGANISM: Homo sapiens
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                                                                                            Local
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CCCGGAAGGCAGTCTGTC 41
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Stolk, John A.
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Lodes, Michael J.
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Smith, Carole L.
                                                                           Conservative
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                                                                                          Score 16.4;
Pred. No. 50;
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RESULT 24 US-09-764-847-1846/c

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Search completed: March Job time: 273 secs
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; ORGANISM: Homo sapiens
US-09-764-847-1846
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US-10-092-154-1845/c
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Best Local Similarity 94.4
Matches 17; Conservative
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SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 1845
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                                                                                                                         Matches
                                                                                                                                                      Query Match
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CURRENT FILING DATE: 2001-01-17
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TITLE OF INVENTION: Nucleic Acids, Proteins,
FILE REFERENCE: PC009
                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Rosen et al.
                                                                                                                                                                                                                     LENGTH: 942
TYPE: DNA
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1: /cgn2_6/ptodata/2/ina/5A_COMB.seq:*
2: /cgn2_6/ptodata/2/ina/5B_COMB.seq:*
3: /cgn2_6/ptodata/2/ina/6A_COMB.seq:*
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Copyright (c) 1993 - 2004 Compugen Ltd.
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US-08-047-041A-12
US-08-047-041A-12
US-08-047-041A-12
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US-08-392-542-1
US-09-147-751-8
US-09-147-751-8
US-09-085-027-1
US-09-085-085-081-084
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(without alignments)
208.107 Million cell updates/sec
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Sequence 15, Appli
Sequence 2, Appli
Sequence 2, Appli
Sequence 4, Appli
Sequence 1, Appli
Sequence 1634, Appli
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 US-09-339-338-81
US-09-643-8263-81
US-09-285-480-81
US-09-285-480-81
US-09-285-480-81
US-09-285-480-81
US-09-285-480-81
US-09-285-9727-12
US-08-445-247-21
US-08-47-179-71
US-08-47-179-71
US-08-47-179-71
US-09-485-247-12
US-09-485-247-12
US-09-485-247-12
US-09-485-247-12
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US-09-103-840A-2
US-09-103-840A-2
US-09-103-840A-2
US-09-103-840A-2
US-09-103-840A-2
US-09-103-847A-2
US-08-256-077-3
US-08-466-127-3
US-08-466-127-3
US-08-256-077-1
US-09-256-93A-35
US-09-256-93A-35
US-09-256-93A-158
US-09-256-93A-158
US-09-256-93A-1524
US-09-256-93A-1544
US-09-256-93A-111544
US-09-256-93A-11154
US-09-256-93A-111544
US-09-256-93A
 Sequence 81, Appl Sequence 81, Appl Sequence 81, Appl Sequence 1, Appl Seq
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Matches

18;

Conservative

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20 CCCGGAAGGCAGTCTGGC 3

CCCGGAAGGCAGTCTGGC

18 0

ALIGNMENTS

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Query Match
Best Local Similarity
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FILING DATE: 06-DEC-1989
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/
AFILING DATE: 29-MAR-1989
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                       NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/928,661
PILLING DATE: 17-AUG-1992
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PC-DOS/MS-DOS
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/047 OF
FILING DATE: 22-Map
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ZIP: 20001.4597
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Kagan, Sarah A.
REGISTRATION NUMBER: 32,141
REFERENCE/DOCKET NUMBER: 011
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                            ORGANISM: HOMO E
                                                                                                                                                                        UBLICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STREET: 1001 G GC
                                                                                                                                                                                                                                                            LECULE TYPE: cDNA POTHETICAL: NO TI-SENSE: NO
                                                                                                                                                                                                                                                                                                                                                                QUENCE CHARACTERISTICS:
LENGTH: 102 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DDRESSEE:
                                                                                                                                                                                             THROMOSOME/SEGMENT:
                                                                                                                                                                                                                                           GINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                 TRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                         ELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   JE OF INVENTION: Detection of Loss of the Wild-Type p53
JE OF INVENTION: Gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           INFORMATION:
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). 5527676
                                                                                                                                                                                                                                                                                                                                               nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              E: Banner & Allegretti, Ltd.
1001 G Street, N.W.
                                                                                       1379-1385
                                                                                                                                                                                                                                                                                                                                                              102 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Fearon, Eric R.
Nigro, Janice M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Vogelstein,
                                                                                                                                         Mol. Cell. Biol
                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                linear
                                                                                                                                                                                                                                                                                                                                  double
 100.0%;
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Score 18;
Pred. No.
                Length 102;
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RESULT 2
US-08-047-041A-2/c
US-08-047-041A-2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION: 435
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: US 07/928,661
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: PatentIn Rel CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ZIP: 20001.4597
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION:
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                                                                                                                                 JBLICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: FILING DATE: 22-MAI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0
                                                                                                                                                                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER: 01
LECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                      REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER:
              PAGES: 245-252
DATE: 1988
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER:
                                                                             ITLE: A variation in the structure of the ITLE: protein-coding region of the human p53
                                                                                                                                                                                                                                                                                                                      MATION FOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                E OF INVENTION:
                                                                                                                                                                                                                                                                                                      ENCE CHARACTERISTICS
                                                                                                                                                                                                 CULE TYPE: '
                                                                                                                                                                                                                                                  RANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                      LEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ING DATE: 06-DEC-1989
APPLICATION DATA:
                                                                                                                                                                   NISM: Homo sapiens
                                                                                                                                                 MOSOME/SEGMENT:
                                                                                                                                                                                                                                                nucleic acid
DEDNESS: double
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                                                                Gene
                                                                                                                                                                                                                                   linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US 07/330,566
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Query Match

100.0%; Score 18;

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Matches

18;

Conservative

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Mismatches

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Indels

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100.0%;

Pred. No. 3.8;

Best Local Similarity

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US-08-795-006A-22/c
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Patent No. 5840579
                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
GENERAL INFORMATION:
APPLICANT: Boeke, Jef
APPLICANT: Brachmann, Rainer
TITLE OF INVENTION: NUCLEIC ACIDS ENCODING P53
TITLE OF INVENTION: MUTATIONS WHICH SUPPRESS P53 CANCER MUTA-TIONS NUMBER OF SEQUENCES: 32
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OPERATING SYSTEM: DOS
SOFTWARE: FRSESEQ for Windows Version 2.
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/795,006A
FILING DATE: 05-FEB-1997
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                              INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BENERAL INFORMATION:

APPLICANT: Boeke, Jef

APPLICANT: Brachmann,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ent No.
                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
LENGTH: 133 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Kagan, Sarah A
REGISTRATION NUMBER: 32141
REFERENCE/DOCKET NUMBER: 01
"ELECOMMUNICATION INFORMATION:
TELEPHONE: 202-508-9100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: NUCLEIC ACTUAL OF INVENTION: MUTATIONS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE:
TTORNEY/AGENT INFORMATION:
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1001 G Street, NW
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            IBM Compatible
                                                                                                                                                                                                                                                                                                                                     linear
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                                                                                                                                                                                                                                                                       100.0%;
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VIEW PERSON P53 CANCER MUTA- TIONS
                                                                                                                                                                                                                                                                       Score 18;
Pred. No. 3
                                                                                                                                                                                                                                                          Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local :
                                                     GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                      APPLICANT: BHATNAGAR, SA
APPLICANT: NAZAKENKO, IX
TITLE OF INVENTION: METH
TITLE OF INVENTION: SEQU
NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: MURAL, RODERT G.
REGISTRATION NUMBER: 28,531
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Kagan, Sarah A
REGISTRATION NUMBER: 32141
REFERENCE/DOCKET NUMBER: 01
TELECOMMINICATION INFORMATION:
TELEPHONE: 202-508-9100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM: MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STREET: 1001 G SC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE: Banner & Witcoff
STREET: 1001 G Street, NW
                                                                                                                                                                                                                                ADDRESSEE: BURNS, DOANE, SWECKER & MATHIS
STREET: P.O. Box 1404
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                              35 CCCGGAAGGCAGTCTGGC 18
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: United States
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                                                                                                                                                                                                                                                                                                                  GEORGE, Jr., Albert
BHATNAGAR, Satish K.
NAZARENKO, Irena
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IBM Compatible
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                                                                                                                                                                                                                                                                                      METHOD FOR ANALYZING A NUCLEOTIDE SEQUENCE AND KIT THEREFOR
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Pred. No. 3.8;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 133;
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RESULT 7
US-08-047-041A-13/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 4
                                                                                                                                                                                                                                                                                                       Sequence 13, Application US/08047041A Patent No. 5527676
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 100.0%; Score 18; DB Best Local Similarity 100.0%; Pred. No. 4.3;
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Best Local &
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Dell'Acqua, Giorgio
APPLICANT: Mann, Michael J.
APPLICANT: Dzau, Victor J.
APPLICANT: Dzau, Victor J.
TITLE OF INVENTION: Compositions and Methods Based Upon an Isoform of p53
FILE REFERENCE: p53f
CURRENT APPLICATION NUMBER: US/09/414,436
CURRENT FILING DATE: 1999-10-07
EARLIER APPLICATION NUMBER: 60/103,849
EARLIER FILING DATE: 1999-10-09
RARLIER FILING DATE: 1999-10-09
RARLIER FILING DATE: 1999-10-09
RARLIER FILING DATE: 1999-10-09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ENGTH: 1070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
LENGTH: 800 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                      APPLICANT: Vogelstein, Bert
APPLICANT: Baker, Suzanne J.
APPLICANT: Fearon, Eric R.
APPLICANT: Nigro, Janice M.
ZIP: 20001.4597
COMPUTER READABLE FORM:
                                                                                                                                   NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REFERENCE/DOCKET NUMBER: 02
TELECOMMUNICATION INFORMATION: (703) 836-6620
                                                                                                                                                   TILE OF INVENTION: Detection OF INVENTION: Gene
                                                                                             STREET:
                                      COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY:
                                                                          CITY: Washington
                                                                                                                 ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                      127 CCCGGAAGGCAGTCTGGC 110
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                                                                                                                                                                                                                                                                                         INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               18;
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                                                        D.C
                                                                                             E: Banner & Allegretti, Ltd
1001 G Street, N.W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
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                                                                                                                                                                                        Detection of Loss of the Wild-Type p53
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US-08-047-041A-12/c
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US-08-047-041A-13
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                                                                                                                                                                                                                                                                                                                                                                                       Query Match 100.0%;
Best Local Similarity 100.0%;
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APPLICATION NUMBER: US 07/446,584
FILING DATE: 06-DEC-1989
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/330,566
APPLICATION NUMBER: US 07/330,566
FILING DATE: 29-MAR-1989
ATTORNEY/AGENT INFORMATION:
NAME: Kagan, Sarah A.
REGISTRATION NUMBER: 32,141
                                                APPLICANT: Vogelstein, Bert
APPLICANT: Baker, Suzanne J.
APPLICANT: Baker, Suzanne J.
APPLICANT: Wigro, Janice M.
TITLE OF INVENTION: Detection
TITLE OF INVENTION: Gene
NUMBER OF SEQUENCES: 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: 202-508-9299
INFORMATION FOR SEQ ID NO: 13:
                                                                                                                                                                            GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                       VOLUME: 6
ISSUE: 12
IAGES: 4650-4656
                                                                                                                                                                                             tent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US/08/047,041/
FILING DATE: 22-MAR-1993
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/928,661
FILING DATE: 17-AUG-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PUBLICATION INFORMATION:
AUTHORS: Harris, N.
JOURNAL: Mol. Cell. Biol
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PAtentIn Release #1.0, Version
CURRENT APPLICATION DATA:
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ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELECOMMUNICATION INFORMATION:
                                     CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MAP POSITION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: nucleic acid
STRANDEDNESS: doub
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COMPUTER: IF
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                  ADDRESSEE:
                                                                                                                                                                                                                                                                                                    113 CCCGGAAGGCAGTCTGGC 96
                                                                                                                                                                                               12, Application US/08047041A
5. 5527676
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E: Banner & Allegretti, Ltd
1001 G Street, N.W.
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                                                                                                                                                                                                                                                                                                                                                                           Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              linear
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22-MAR-1993
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                                                                                        Detection of Loss of the Wild-Type p53
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Pred. No. 4
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: Washington E: D.C. TRY: USA

COMPUTER READABLE FORM: MEDIUM TYPE: Floppy

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS

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                                                                                                                                                             Sequence 1, Application US/07912011 Patent No. 5382510
                                                                                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: 202-508-9299
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
LENGTH: 1307 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ANTI-SENSE: NO
ORIGINAL SOURCE:
ORGANISM: Homo sapiens
POSITION IN GENOME:
MAP POSITION: 17p13.1
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MOLECULE TYPE: CI
HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: US 07/446,584
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION DATA:
PRIOR APPLICATION NUMBER: US 07/928,661
PRIOR DATE: 17-AUG-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US 07/446,584
FILING DATE: 06-DEC-1989
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/330,566
                                                                                            APPLICANT: Levine, Arnold J
APPLICANT: Shenk, Thomas E.
APPLICANT: Finlay, Cathy A.
               NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
ADDRESSEE: Inclone Systems Incorporated
STREET: 180 Varick Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PUBLICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Kagan, Sarah A.
REGISTRATION NUMBER: 32,141
REFERENCE/DOCKET NUMBER: 01
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: Patentin Release #1.0, Version #1.25
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ATTORNEY/AGENT INFORMATION:
                                                                            TLE OF INVENTION: Probes for Detecting Mutant p53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: FILING DATE: 22-MAI CLASSIFICATION: 43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  JOURNAL:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AUTHORS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TIE: human p53 protein
URNAL: Mol. Cell. Biol.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               THORS: Harris, N.
TLE: Molecular basis for heterogeneity of the
                                                                                                                                                                                                                                                            117
                                                                                                                                                                                                                                                                                                                          18;
                                                                                                                                               INFORMATION:
                                                                                                                                                                                                                                                                                       1 CCCGGAAGGCAGTCTGGC 18
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New York
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                                                                                                                                Arnold J.
                                                                                                                                                                                                                                                                                                                                    100.0%; Score 18; 100.0%; Pred. No.
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TELECOMMUNICATION OF TELEPANE: 212-045-2054
TELEPAX: 212-045-2054
INFORMATION FOR SEQ 1D NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 1317 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TYPE: TYPE: Tiple: Tiple: TYPE: Tiple: Tiple: Tiple: TYPE: Tiple: Tiple: TYPE: Tiple: Tip
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Best Local Similarity
Matches 18; Conserva
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US-08-347-792-1/c
                                                                                                                   ATFLIANCE DATE:
FILING DATE:
CLASSIFICATION: 530
ATTORNEY/AGENT INFORMATION:
NAME: Bak, Mary E.
REGISTRATION NUMBER: 31,215
REFERENCE/DOCKET NUMBER: WST5
TELECOMMUNICATION INFORMATION:
TELEPHONE: 215-540-9206
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                sequence 1, Application US/08347792
Patent No. 5573925
                                                                           TELEFAX: 215-540-5818
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
NAME: Feit, Irving N.
REGISTRATION NUMBER: 28,6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/912,011
FILING DATE: 10-JUL-1992
CLASSIFICATION: 435
SEQUENCE CHARACTERISTICS:
LENGTH: 1317 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Halazonetis, Thanos D.
TITLE OF INVENTION: p53 Proteins With Altered
TITLE OF INVENTION: Tetramerization Domains
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDIUM TYPE: Floppy
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LECOMMUNICATION INFORMATION:
TELEPHONE: 212-645-1405
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE: Howson and Howson
STREET: Spring House Corporate Cntr.,
CITY: Spring House
STATE: Pennsylvania
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/08/347,792
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IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
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100.0%; Pred. No.
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US-08-347-792-1
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В
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; LOCATION:
US-08-431-357-1
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Patent No. 5721340
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                                                                                  Query Match
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                                                                                                                                                                                                                                                             TELEFAX: 215-540-5818
INFORMATION FOR SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION: 435
PRIOR APPLICATION UNMBER: US 08/347,792
PILLING DATE: 28-NOV-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patenth Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/431,357
FILING DATE:
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STREET: Spring Hour
CITY: Spring Hour
Thire: Pennsylvania
TISA
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LENGTH: 1317 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Halazonetis, Thanos D.
TITLE OF INVENTION: p53 Proteins With Altered
TITLE OF INVENTION: Tetramerization Domains
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES: 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: 1:
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                                                                      Local
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                                                                                                                                                                                          STRANDEDNESS:
TOPOLOGY: 111
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Similarity 100.0%;
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CCCGGAAGGCAGTCTGGC 110
                          ccceenaeccaercreec 18
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215-540-5818
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                                                     100.0%; S
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Pred. No.
                                                                      Score 18;
Pred. No.
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US-08-697-221-1/c
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Patent No. 5847083
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEPHONE: 215-540-920
TELEFAX: 215-540-5818
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                equence 1, Application US/08392542 atent No. 6169073 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REFERENCE/DOCKET NUMBER: WS
TELECOMMUNICATION INFORMATION:
TELEPHONE: 215-540-9206
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION:
NAME: Kodroff, Cathy A.
REGISTRATION NUMBER: 33,980
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 6
FILING DATE: 22-SEP-1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Halazonetis, Thanos D.
TITLE OF INVENTION: Modified p53 Constructs and Uses
TITLE OF INVENTION: Therefor
NUMBER OF SEQUENCES: 33
                                                               TITLE OF INVENTION: FUNDMBER OF SEQUENCES: 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MOLECULE TYPE:
                                                                                                                                                   APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME/KEY:
LOCATION:
                                                                                                                 PPLICANT: Hartwig, Wolfgang
ITLE OF INVENTION: Peptides nad Peptidomimetics with
ITLE OF INVENTION: Structural Similarity to Human p53 That Activate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 1317 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
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                                                   DDRESSEE:
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                Washington,
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District of Columbia
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                             Banner, Birch, McKie
1001 G Street, N.W.
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136..1314
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Pred. No.
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OMPUTER READABLE FORM: MEDIUM TYPE: Floppy

Floppy disk

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Query Match
Best Local Similarity
Watches 18; Conserva
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US-08-894-327-1/c
RESULT 15
US-09-147-751-8/c
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SEQ ID NO 1
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Patent No. 6245886
GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 1317 base pairs
                                                                                                                                                                                                                                                                                                                                                  EARLIER APPLICATION NUMBER: pctus96/01535
EARLIER FILING DATE: 1996-02-16
EARLIER APPLICATION NUMBER: 08/392,542
EARLIER FILING DATE: 1995-02-16
                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQ ID NOS: 35
                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/08/894,327
CURRENT FILING DATE: 1997-12-04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Halazonetis,
APPLICANT: Hartwig, Wol
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NAME: POBOTSKe, Laurence H.
REGISTRATION NUMBER: 34,698
REFERENCE/DOCKET NUMBER: 0486
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202 508-9100
                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILE REFERENCE: 2973.19998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RPPLICANT: Hartwig, Wolfgang
NITLE OF INVENTION: Peptides and peptidomimetics with
NITLE OF INVENTION: structural similarity to human p53 that activate p53
NITLE OF INVENTION: function
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ORIGINAL SOURCE:
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                            ENGTH: 1317
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EDNESS: double
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                                                                                                                                                               100.0%; Score 18; DB 3; 100.0%; Pred. No. 4.3;
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Pred. No.
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                                                                                                                                                                                      RESULT 16
                                                                                                             GENERAL INFORMATION:
                                                                                                                           Sequence 1, Application US/09305914
Patent No. 6388062
                                                                                                                                                                                                                                                                                                                       Matches
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                  APPLICANT: Halazonetis, Thanos D.
APPLICANT: Stavridi, Elena S.
TITLE OF INVENTION: Modified p53 Tetramerization Domains Having Hydrophobic
TITLE OF INVENTION: Amino Acid Substitutions
FILE REFERENCE: WST84AUSA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: (202) 672-5399 INFORMATION FOR SEQ ID NO:
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CURRENT APPLICATION NUMBER: US/09/305,914
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TITLE OF INVENTION: N
TITLE OF INVENTION: U
TITLE OF INVENTION: U
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
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ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER: WO PCT/JP97/03019
FILING DATE: 29-AUG-1997
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
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TOPOLOGY: lir
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Maebius, Stephen B.
REGISTRATION NUMBER: 35,264
REFERENCE/DOCKET NUMBER: 84
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: JP 8/347090 FILING DATE: 26-DEC-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: 18 CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                          Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: JP 8/229061 FILING DATE: 29-AUG-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US/0
FILING DATE: 18-MAY-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPHONE:
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                                                                                                                                                                                                                                                                                                                                          Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                     nucleic acid
BEDNESS: double
                                                                                                                                                                                                                                                                                   CCCGGAAGGCAGTCTGGC 18
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OBATA, Kazuaki
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KUSUMI, Kayo
                                                                                                                                                                                                                                                                                                                         Conservative
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A, Mikayo
                                                                                                                                                                                                                                                                                                                                        100.0%; Score 18; DB 4; Length 1317; 100.0%; Pred. No. 4.3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DETECTING AND/OR ISOLATING USING RECA-LIKE RECOMBINASE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          672-5300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              METHODS FOR TARGETING, ENRICHING, DETECTING AND/OR ISOLATING TARGET
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ORGANISM: Homo sapiens SEQUENCE DESCRIPTION: SEQ ID NO: 1: US-09-685-027-1
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US-09-685-027-1/c
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Patent No. 6420118
GENERAL INFORMATION:
Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT FILING DATE: 1999-05-05
EARLIER APPLICATION NUMBER: 60/084,839
EARLIER FILING DATE: 1998-05-08
NUMBER OF SEQ ID NOS: 4
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                   TELEFAX: 202 508-9299 INFORMATION FOR SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FEATURE: NAME/KEY: CDS
LOCATION: (136)..(1314)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: DNA
ORGANISM: human p53
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MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.25
                                                                                                                                                   SEQUENCE CHARACTERISTICS:
LENGTH: 1317 base pairs
TYPE: nucleic acid
STRANDENNESS: double
                                                                                                                                                                                                                                                                     NAME: POSOTSKE, Laurence H.
RECISTRATION NUMBER: 34,698
REFERENCE/DOCKET NUMBER: 0486.48439
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202 508-9100
                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: 08/392,542 FILING DATE: <Unknown> ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/685,027
FILING DATE: 10-Oct-2000
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Halazonetis, Thanos
Hartwig, Wolfgang
TITLE OF INVENTION: Peptides nad Peptidomimetics with
                                                                                              ORIGINAL SOURCE:
                                                                                                                 TOPOLOGY: linear MOLECULE TYPE: cDNA
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18; Conservative
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STATE: District of Columbia
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STREET: 1001 G Street, N.W.
CITY: Washington, D.C.
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100.0%; Score 18;
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BB
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Length 1317;
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Matches 18; Conserv
                                                          Matches
                                                                                        Query Match
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INFORMATION FOR SEQ ID NO: 1:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM FOC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE: 01-JUN-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/347,792
FILING DATE: 28-NOV-1994
                                                                                                                                                                   FEATURE:
                                                                                                                                                                             MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                        REFERENCE/DOCKET NUMBER: WS
TELECOMMUNICATION INFORMATION:
TELEPHONE: 215-540-9206
                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
APPLICATION NUMBER: 1
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APPLICANT: and Biology
APPLICANT: Halazonetis, Thanos D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ITLE OF INVENTION: p53 Proteins William OF INVENTION: Tetramerization
                                                                                                                                    NAME/KEY:
LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: PCT/US95/15353
PILING DATE:
CLASSIFICATION:
                                                                           Local
                                                                                                                                                                                                  TOPOLOGY:
                                                                                                                                                                                                            STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                  REGISTRATION NUMBER: 31,215
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 127
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                             1 CCCGGAAGGCAGTCTGGC 18
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18; Conserv
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T: Spring House Corporate Cntr.,
Spring House
: Pennsylvania
                                                                                                                                                                                                                              nucleic acid
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CCCGGAAGGCAGTCTGGC 110
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                                                          Conservative
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136..1314
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                                                                       100.0%;
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                                                                           Score 18;
Pred. No.
                                                            Mismatches
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                                                                                         Length 1317;
                                                            Indels
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RESULT 19 US-09-404-879A-332

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RESULT 20
US-09-702-705-1634/c
; Sequence 1634, Application US/09702705
natent No. 6504010
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                                                                                                                                                                                                                                                                                                  ; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-702-705-1634
                                                                                                           US-09-736-457-1634/c
                                                                                                                             RESULT 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: King, Gordon E.
APPLICANT: Algate, Paul A.
APPLICANT: Algate, Paul A.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.462C2
CURRENT APPLICATION NUMBER: US/09/404,879A
CURRENT FILING DATE: 1999-09-24
NUMBER OF SEQ ID NOS: 393
SOFTWARE: FASTSEQ for Windows Version 3.0
SEQ ID NO 332
LENGTH: 184
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Bangur Chaitanya S
APPLICANT: Lodes, Michael A.
APPLICANT: Fanger, Gary
APPLICANT: Vedvick, Tom
APPLICANT: Carter, Darrick
APPLICANT: Retter, Marc
APPLICANT: Mannion, Jane
APPLICANT: Fan, Liqun
                                                                                                                                                                                                                           Query Match
Best Local Similarity 94.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 332, Application US/09404879A Patent No. 6468546 GENERAL INFORMATION:
                                                                              Sequence 1634, Application US/09736457 Patent No. 6509448
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Best Local :
                                                                 GENERAL INFORMATION:
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                APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND TITLE OF INVENTION: DIAGNOSIS OF LUNG CANCER FILE REFERENCE: 210121.478C194 CURRENT APPLICATION NUMBER: US/09/702,705 CURRENT FILING DATE: 2000-10-30 NUMBER OF SEQ ID NOS: 1833
                                                                                                                                                                                                                                                                                                                                                   ENGTH: 447
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Local Similarity 94.1%;
nes 16; Conservative
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Wang, Tongtong
Bangur, Chaitanya S.
Lodes, Michael A.
Fanger, Gary
                                                                                                                                                                            CCAGAAGGCAGTCTGGC 57
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Chaitanya S.
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94.1%;
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Pred. No. 74;
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Pred. No. 78
                                                                                                                                                                                                                                        Mismatches
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US-09-671-325-1634/c
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US-09-614-124B-1634/c
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Matches 16
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Matches 16
                                                                                           Sequence 1634, Application US/09671325
Patent No. 6667154
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                                                                                INFORMATION:
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Wang, Tongtong
Bangur, Chaitanya S
Lodes, Michael A.
Fanger, Gary
Vedvick, Tom
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; TYPE: DNA; Homo sapiens US-09-614-124B-1634
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Fan, Liquin APPLICANT: Wang, Aijun APPLICANT: Wang, Aijun APPLICANT: Wang, Aijun COMPOSITIONS AND METHODS FOR THE THERAPY AND TITLE OF INVENTION: DIAGNOSIS OF LUNG CANCER FILE REFERENCE: 210121.478C15
CURRENT APPLICATION NUMBER: US/09/736,457
CURRENT FILING DATE: 2000-12-13
NUMBER OF SEQ ID NOS: 1864
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 1634
                                                                                                                                                                                                          SOFTWARE: FastSEQ for Windows Version 3.0 SEQ ID NO 1634 LENGTH: 447
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GENERAL INFORMATION:
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CURRENT APPLICATION NUMBER: US/09/614,124B
CURRENT FILING DATE: 2001-07-11
NUMBER OF SEQ ID NOS: 1668
                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Wang, Tongtong
APPLICANT: Bangur, Chaitanya S.
APPLICANT: Lodes, Michael A.
APPLICANT: Fanger, Gary
                                                                                                                                                                                                                                                                                                           PPLICANT: Vedvick, Tom

PPLICANT: Carter, Darrick

PPLICANT: Retter, Marc

PPLICANT: Retter, Marc

PPLICANT: Repter, Marc

PPLICANT: Repter, Marc

PPLICANT: Repter, Marc

PPLICANT: Repter Marc

ITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY AND

ITLE OF INVENTION: DIAGNOSIS OF LUNG CANCER

ILE REFERENCE: 210121.478C9
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les 16; Conservative
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16; Conserv
                              2 CCGGAAGGCAGTCTGGC 18
CCAGAAGGCAGTCTGGC
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Retter, Marc
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                                                                          Conservative
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Pred. No. 78;
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Pred. No. 78;
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Retter, Marc

Carter, Darrick

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; LENGTH: 633
; TYPE: DNA
; ORGANISM: Homo sapiens
; PEATURE:
; NAME/KEY: misc_feature
; LOCATION: (1)...(633)
; OTHER INFORMATION: n = A,T,C or G
US-09-385-982-206
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US-09-385-982-206/c
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US-09-222-575-81
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; ORGANISM: Homo sapiens
US-09-671-325-1634
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CURRENT FILING DATE: 1999-08-30
EARLIER APPLICATION NUMBER: 09/328,111
EARLIER FILING DATE: 1999-06-08
EARLIER APPLICATION NUMBER: 60/117,393
EARLIER FILING DATE: 1999-01-27
EARLIER FILING DATE: 1999-01-27
EARLIER APPLICATION NUMBER: 60/098,639
EARLIER APPLICATION NUMBER: 60/098,639
EARLIER FILING DATE: 1988-08-31
NUMBER OF SEQ ID NOS: 544
SOFTWARE: FastSEQ for Windows Version 3.0
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SEQ ID NO 1634
LENGTH: 447
                                                                         Sequence 81, Application US/09222575
Patent No. 6387697
GENERAL INFORMATION:
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Best Local Similarity
Matches 16; Conserv
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Patent No. 6262334
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQ ID NO 206
APPLICANT: Yuqiu, Jiang
APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Xu, Jiangchun
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/09/671,325
CURRENT FILING DATE: 2000-09-26
NUMBER OF SEQ ID NOS: 1825
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: ENDEGE, WILSON O., ET AL.

TITLE OF INVENTION: NOVEL HUMAN GENES AND GENE EXPRESSION

FIGURE OF INVENTION: PRODUCTS: II
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APPLICANT: Fan, Liqun
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF LUNG CANCER
FILE REFERENCE: 210121.478C12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILE REFERENCE: CCDNA-260XX
                                                                                                                                                                                                                     426
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94.1%; Pred. No. 79;
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                              S
                                                                                                                                   ; TYPE: DNA
; ORGANISM: Human
US-09-222-575-81
                                                                                                                                                                                TITLE OF INVENTION: Compositions for the Treatment and Diagnosis of Breast Cancer TITLE OF INVENTION: and Methods for Their Use FILE REFERENCE: 210121.470
CURRENT APPLICATION NUMBER: US/09/222,575
CURRENT FILING DATE: 1998-12-28
NUMBER OF SEQ ID NOS: 174
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 81
LENGTH: 647
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                                                              / Match 85.6%;
Local Similarity 94.1%;
les 16; Conservative
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                                  2 CCGGAAGGCAGTCTGGC
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AA358056 BG608755 CH270401 RE1295366 RE25637 CG778171 BF706710 BF706710 BF706710 CG77814972 CG344533 CG7782144 CG7782144 CG7782144 CG7782158 CG782158 CG7821666 CG7821666 CG78216666 CG78216666 CG782166666 CG7821666666666666666666666666666666666666	3573 8516 54672	26780 3978 15695	29906	.9229 1081	35038	53938 6211	144765 36739 113221	98893 86919	19531 14676 7581	8353 35676	00000 000000	9965	9631	94801	44316	43070	51887	8090	96796 02L60	1861 5141	1702	45594	38352 66114 00862	57354
AA358056 EST766889 BG608755 307436 MA CB270401 1009308 H BI295366 UI-R-DK0- R25637 yh45h08.11 CG778171 1123026E1 BF706710 281330 MA BG814972 dab98c10. CG344533 OG6AB0OTC CB776194 AMGNNUC:S CG7782144 112304BG0 CG7783186 1123016E1 CG782148 112304BG0 CG778216 112304BG0 CG7783186 112304BG1 CC8802 CR802 Rice BB68016 1B680161 BM482266 535083 MA AQ006068 CIT-HSP-2 BM106200 509869 MA AA329660 EST33298	AI93357 AA32851 BY3546	AA33025 AW3267 AA33397 BI0156	CB0299	AV01922 AA33108	BX3350	AZ3539 AL03621	BM5447 BQ4367 BM0132	BG6988 BE7869	BQ9195 BI1146	AU11835 BG4356	BF3160	AU11996	BG6215 AU13963	BG2948	BE1443	BF0430	BF6518	AA29809	BX3967	AA17186 AU13514	BM4692	BX3455	BG3383 BM4661 BE9008	BU1573
56 EST66 1009 101 1009 106 UT F 7 Yh4510 77 11123 77 1123 77 1281 77 0 281 77 0 281 77 0 281 78 0 406 33 0 66 33 0 66 33 0 66 33 1 112 3 84 112 3 88 112 3 58 112 5 8 112 5 8 15 5 8 5 5 8	73 wn40f 6 EST32 72 BY35	52 EST33 780 1977 78 EST38 595 PMO-	06 TGES	9 AV019	175 6018)38 BX33	38 1MOC	765 AGEN 739 AGEN 221 6036	393 6027 319 6014	31 AGEN 576 6028	3 AU118	00 AGEN	55 AU119	1 AU139	01 6023	16 MR0-)70 BP25	387 2747	0 EST11	196 BX39 125 Teti	11 ZD22c)2 AL521	77 AL530	152 6024 114 AGEN 362 6016	54 AGEN
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RESULT 1 AU076984 LOCUS LOCUS DEFINITI ACCESSIO VERSION VERSION KEYWORDS SOURCE ORGANI ORGANI REFERENC AUTHOR		NUL	c 196		191 c 192 c 193	c 189			C 181		C 178			0 171	169	C 167	0 100	c 163		c 159			c 152 c 153	_
SULT 1 075984/c CUS FINITION FINITION RSION RSION YWORDS URCE ORGANISM ORGANISM FERRENCE AUTHORS		198 199 1 200 1	196 1	194 1 195 1	192 1	189 1		184	181				173 . 1	ے ب	, ب	, بر د	7 17 1	63 1	161 1 162 1	159 1	158 1	1000		151 1
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AU076984 AU076984 similar t similar t mRNA sequ MRNA sequ AU076984 AU076984 AU076984 EST. bomo sap: Eukaryott Mammalia Li (basea Suzuki, y TSUNOda, T		198 15.4 85.6 199 15.4 85.6 200 15.4 85.6	196 15.4 85.6 197 15.4 85.6	194 15.4 85.6 195 15.4 85.6	191 15.4 85.6 192 15.4 85.6	189 15.4 85.6 190 15.4 85.6	186 15.4 85.6 187 15.4 85.6 188 15.4 85.6	184 15.4 85.6 185 15.4 85.6	181 15.4 85.6 182 15.4 85.6	179 15.4 85.6 180 15.4 85.6	177 15.4 85.6	175 15.4 85.6	173 , 15.4 85.6	15.4	15.4 85.6	15.4 85.6	55 15.4 85.6 15.4 85.6	63 15.4 85.6	161 15.4 85.6 162 15.4 85.6	159 15.4 85.6 160 15.4 85.6	158 15.4 85.6	15.4 85.6	152 15.4 85. 153 15.4 85. 154 15.4 85.	151 15.4 85.6
AU076984 AU076984 Sugano cDN similar to 5'-end r mRNA sequence. AU076984 AU076984.1 GI:7439 EST. Homo sapiens (human Homo sapiens Eukaryota; Metazoa; Mammalia; Eutheria; 1 (bases 1 to 190) Suzuki, Y., Ishihara and Sugano, S. Statistical analysi		198 15.4 85.6 570 12 199 15.4 85.6 571 13 200 15.4 85.6 571 28	196 15.4 85.6 562 14 197 15.4 85.6 564 9	194 15.4 85.6 556 10 195 15.4 85.6 559 29	191 15.4 85.6 545 9 192 15.4 85.6 550 9 193 15.4 85.6 555 10	189 15.4 85.6 539 14 190 15.4 85.6 540 12	186 15.4 85.6 521 10 187 15.4 85.6 524 12 188 15.4 85.6 530 12	184 15.4 85.6 515 12 185 15.4 85.6 518 28	181 15.4 85.6 510 14 182 15.4 85.6 511 12	179 15.4 85.6 509 10 180 15.4 85.6 510 10	178 15.4 85.6 506 9 178 15.4 85.6 506 9	174 15.4 85.6 504 10 175 15.4 85.6 504 10	173 , 15.4 85.6 497 10	15.4 85.6 495 10	15.4 85.6 485 12	15.4 85.6 483 10	65 15.4 85.6 478 14 66 15.4 85.6 480 14	63 15.4 85.6 478 12	161 15.4 85.6 465 10 162 15.4 85.6 470 10	159 15.4 85.6 452 9 160 15.4 85.6 454 13	158 15.4 85.6 440 12	155 15.4 85.6 428 14 156 15.4 85.6 436 9	152 15.4 85.6 424 14 153 15.4 85.6 427 14 154 15.4 85.6 428 9	151 15.4 85.6 424 12
AU076984 AU076984 Sugano cDNA lib similar to 5'-end region mRNA sequence. AU076984 AU076984 I GI:7439546 EST. Homo sapiens (human) Homo sapiens Eukaryota; Metazoa; Chor Mammalia; Eutheria; Prim 1 (bases 1 to 190) Suzuki, Y., Ishihara, D., Tsunoda, T., Watanabe, M., and Sugano, S. Statistical analysis of	ALIGN	198 15.4 85.6 570 12 199 15.4 85.6 571 13 200 15.4 85.6 571 28	196 15.4 85.6 562 14 197 15.4 85.6 564 9	194 15.4 85.6 556 10 195 15.4 85.6 559 29	191 15.4 85.6 545 9 192 15.4 85.6 550 9 193 15.4 85.6 555 10	189 15.4 85.6 539 14 190 15.4 85.6 540 12	186 15.4 85.6 521 10 187 15.4 85.6 524 12 188 15.4 85.6 530 12	184 15.4 85.6 515 12 185 15.4 85.6 518 28	181 15.4 85.6 510 14 182 15.4 85.6 511 12	179 15.4 85.6 509 10 180 15.4 85.6 510 10	178 15.4 85.6 506 9 178 15.4 85.6 506 9	174 15.4 85.6 504 10 175 15.4 85.6 504 10	173 , 15.4 85.6 497 10	15.4 85.6 495 10	15.4 85.6 485 12	15.4 85.6 483 10	65 15.4 85.6 478 14 66 15.4 85.6 480 14	63 15.4 85.6 478 12	161 15.4 85.6 465 10 162 15.4 85.6 470 10	159 15.4 85.6 452 9 ALO36	158 15.4 85.6 439 10 BF11	155 15.4 85.6 428 14 R586 156 15.4 85.6 436 9 N1758	152 15.4 85.6 424 14 W302 153 15.4 85.6 427 14 CF11 154 15.4 85.6 428 9 A1909	151 15.4 85.6 424 12 BI8173
AU076984 AU076984 Sugano cDNA library Homo similar to 5'-end region of Human mRNA sequence. AU076984 AU076984 I GI:7439546 EST. Homo sapiens (human) Homo sapiens Eukaryota; Metazoa; Chordata; Crar Mammalia; Eutheria; Primates; Cata 1 (bases 1 to 190) Suzuki, Y., Ishihara, D., Sasaki, M., Tsunoda, T., Watanabe, M., Komatsu, T and Sugano, S. Statistical analysis of the 5' unt	ALIGNMENTS	198 15.4 85.6 570 12 199 15.4 85.6 571 13 200 15.4 85.6 571 28	196 15.4 85.6 562 14 197 15.4 85.6 564 9	194 15.4 85.6 556 10 195 15.4 85.6 559 29	191 15.4 85.6 545 9 192 15.4 85.6 550 9 193 15.4 85.6 555 10	189 15.4 85.6 539 14 190 15.4 85.6 540 12	186 15.4 85.6 521 187 15.4 85.6 524 188 15.4 85.6 530	184 15.4 85.6 515 12 185 15.4 85.6 518 28	181 15.4 85.6 510 14 182 15.4 85.6 511 12	179 15.4 85.6 509 10 180 15.4 85.6 510 10	178 15.4 85.6 506 9 178 15.4 85.6 506 9	174 15.4 85.6 504 10 175 15.4 85.6 504 10	173 , 15.4 85.6 497 10	15.4 85.6 495 10	15.4 85.6 485 12	15.4 85.6 483 10	65 15.4 85.6 478 14 66 15.4 85.6 480 14	63 15.4 85.6 478 12	161 15.4 85.6 465 10 162 15.4 85.6 470 10	159 15.4 85.6 452 9 ALO36	158 15.4 85.6 439 10 BF11	155 15.4 85.6 428 14 156 15.4 85.6 436 9	152 15.4 85.6 424 14 W302 153 15.4 85.6 427 14 CF11 154 15.4 85.6 428 9 A1909	151 15.4 85.6 424 12 BI8173
AU076984 AU076984 Sugano cDNA library Homo similar to 5'-end region of Human mRNA sequence. AU076984 AU076984 I GI:7439546 EST. Homo sapiens (human) Homo sapiens Eukaryota; Metazoa; Chordata; Crar Mammalia; Eutheria; Primates; Cata 1 (bases 1 to 190) Suzuki, Y., Ishihara, D., Sasaki, M., Tsunoda, T., Watanabe, M., Komatsu, T and Sugano, S. Statistical analysis of the 5' unt	ALIGNMENTS	198 15.4 85.6 570 12 199 15.4 85.6 571 13 200 15.4 85.6 571 28	196 15.4 85.6 562 14 197 15.4 85.6 564 9	194 15.4 85.6 556 10 195 15.4 85.6 559 29	191 15.4 85.6 545 9 192 15.4 85.6 550 9 193 15.4 85.6 555 10	189 15.4 85.6 539 14 190 15.4 85.6 540 12	186 15.4 85.6 521 10 187 15.4 85.6 524 12 188 15.4 85.6 530 12	184 15.4 85.6 515 12 185 15.4 85.6 518 28	181 15.4 85.6 510 14 182 15.4 85.6 511 12	179 15.4 85.6 509 10 180 15.4 85.6 510 10	178 15.4 85.6 506 9 178 15.4 85.6 506 9	174 15.4 85.6 504 10 175 15.4 85.6 504 10	173 , 15.4 85.6 497 10	15.4 85.6 495 10	15.4 85.6 485 12	15.4 85.6 483 10	65 15.4 85.6 478 14 66 15.4 85.6 480 14	63 15.4 85.6 478 12	161 15.4 85.6 465 10 162 15.4 85.6 470 10	159 15.4 85.6 452 9 ALO36	158 15.4 85.6 439 10 BF11	155 15.4 85.6 428 14 R586 156 15.4 85.6 436 9 N1758	152 15.4 85.6 424 14 W302 153 15.4 85.6 427 14 CF11 154 15.4 85.6 428 9 A1909	151 15.4 85.6 424 12 BI8173
AU076984 AU076984 Sugano cDNA library Homo similar to 5'-end region of Human mRNA sequence. AU076984 AU076984 I GI:7439546 EST. Homo sapiens (human) Homo sapiens Eukaryota; Metazoa; Chordata; Crar Mammalia; Eutheria; Primates; Cata 1 (bases 1 to 190) Suzuki, Y., Ishihara, D., Sasaki, M., Tsunoda, T., Watanabe, M., Komatsu, T and Sugano, S. Statistical analysis of the 5' unt	ALIGNMENTS	198 15.4 85.6 570 12 BG530001 199 15.4 85.6 571 13 BQ126161 200 15.4 85.6 571 28 AZ109510	196 15.4 85.6 562 14 CB266565 197 15.4 85.6 564 9 AUZ80282	194 15.4 85.6 556 10 AW388278 195 15.4 85.6 559 29 CC753990	191 15.4 85.6 545 9 A11/7319 192 15.4 85.6 550 9 AA277412 193 15.4 85.6 555 10 BE634376	189 15.4 85.6 539 14 CB265876 190 15.4 85.6 540 12 BG537072	186 15.4 85.6 521 10 AW838191 187 15.4 85.6 524 12 BI817305 188 15.4 85.6 530 12 BM721923	184 15.4 85.6 515 12 BI775888 185 15.4 85.6 518 28 AZ296219	181 15.4 85.6 510 14 CB143418 182 15.4 85.6 511 12 B1817644 183 15.4 85.6 514 0 13184037	179 15.4 85.6 509 10 AW673601 180 15.4 85.6 510 10 AW520875	177 15.4 85.6 506 14 CB723596	174 15.4 85.6 504 10 BE937951 175 15.4 85.6 504 10 BE833812 176 15.4 85.6 505 14 OR569000	173 , 15.4 85.6 497 9 AALLI/07	15.4 85.6 495 10 AV815788	15.4 85.6 485 12 BJ061283	15.4 85.6 483 10 AW630245	65 15.4 85.6 478 14 W49749	63 15.4 85.6 478 12 BG961196	161 15.4 85.6 465 10 AW885823 162 15.4 85.6 470 10 BE106945	159 15.4 85.6 452 9 ALO36065 160 15.4 85.6 454 13 BY482752	157 15.4 85.6 439 10 BF117509	155 15.4 85.6 428 14 R58670 156 15.4 85.6 436 9 R1759475	15.2 15.4 85.6 424 14 W30204 15.3 15.4 85.6 427 14 CF112695 15.4 15.4 85.6 428 9 A1909705	151 15.4 85.6 424 12 BI817349
AU076984 Sugano cDNA library Homo sapiens cDNA clone similar to 5'-end region of Human p53 cellular tumor mRNA sequence. AU076984 Sugano cDNA library Homo sapiens cDNA clone similar to 5'-end region of Human p53 cellular tumor mRNA sequence. AU076984 AU076984.1 GI:7439546 EST. Homo sapiens (human) Homo sapiens (human) Homo sapiens Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; 1 (bases 1 to 190) Suzuki, Y., Ishihara, D., Sasaki, M., Nakagawa, H., Hata Tsunoda, T., Watanabe, M., Komatsu, T., Ota, T., Isogai, and Sugano, S. Statistical analysis of the 5' untranslated region o	ALIGNMENTS	198 15.4 85.6 570 12 BG530001 199 15.4 85.6 571 13 BQ126161 200 15.4 85.6 571 28 AZ109510	196 15.4 85.6 562 14 CB266565 197 15.4 85.6 564 9 AUZ80282	194 15.4 85.6 556 10 AW388278 195 15.4 85.6 559 29 CC753990	191 15.4 85.6 545 9 A11/7319 192 15.4 85.6 550 9 AA277412 193 15.4 85.6 555 10 BE634376	189 15.4 85.6 539 14 CB265876 190 15.4 85.6 540 12 BG537072	186 15.4 85.6 521 10 AW838191 187 15.4 85.6 524 12 BI817305 188 15.4 85.6 530 12 BM721923	184 15.4 85.6 515 12 BI775888 185 15.4 85.6 518 28 AZ296219	181 15.4 85.6 510 14 CB143418 182 15.4 85.6 511 12 B1817644 183 15.4 85.6 514 0 13184037	179 15.4 85.6 509 10 AW673601 180 15.4 85.6 510 10 AW520875	177 15.4 85.6 506 14 CB723596	174 15.4 85.6 504 10 BE937951 175 15.4 85.6 504 10 BE833812 176 15.4 85.6 505 14 OR569000	173 , 15.4 85.6 497 9 AALLI/07	15.4 85.6 495 10 AV815788	15.4 85.6 485 12 BJ061283	15.4 85.6 483 10 AW630245	65 15.4 85.6 478 14 W49749	63 15.4 85.6 478 12 BG961196	161 15.4 85.6 465 10 AW885823 162 15.4 85.6 470 10 BE106945	159 15.4 85.6 452 9 ALO36065 160 15.4 85.6 454 13 BY482752	157 15.4 85.6 439 10 BF117509	155 15.4 85.6 428 14 R58670 156 15.4 85.6 436 9 R1759475	15.2 15.4 85.6 424 14 W30204 15.3 15.4 85.6 427 14 CF112695 15.4 15.4 85.6 428 9 A1909705	151 15.4 85.6 424 12 BI817349
AU076984 Sugano cDNA library Homo sapiens cDNA clone similar to 5'-end region of Human p53 cellular tumor mRNA sequence. AU076984 AU076984 GI:7439546 EST. Homo sapiens (human) Homo sapiens (chordata; Craniata; Vertebrata; I man sapiens Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; I chase; I to 190) Suzuki, Y., Ishihara, D., Sasaki, M., Nakagawa, H., Hata, Tsunoda, T., Watanabe, M., Komatsu, T., Ota, T., Isogai, Tand Sugano, S. Statistical analysis of the 5' untranslated region of Statistical analysis of the Statistical analysis of the 5' untranslated region of Statist	ALIGNMENTS	198 15.4 85.6 570 12 199 15.4 85.6 571 13 200 15.4 85.6 571 28	196 15.4 85.6 562 14 CB266565 197 15.4 85.6 564 9 AUZ80282	194 15.4 85.6 556 10 AW388278 195 15.4 85.6 559 29 CC753990	191 15.4 85.6 545 9 A11/7319 192 15.4 85.6 550 9 AA277412 193 15.4 85.6 555 10 BE634376	189 15.4 85.6 539 14 CB265876 190 15.4 85.6 540 12 BG537072	186 15.4 85.6 521 10 AW838191 187 15.4 85.6 524 12 BI817305 188 15.4 85.6 530 12 BM721923	184 15.4 85.6 515 12 BI775888 185 15.4 85.6 518 28 AZ296219	181 15.4 85.6 510 14 CB143418 182 15.4 85.6 511 12 B1817644 183 15.4 85.6 514 0 13184037	179 15.4 85.6 509 10 AW673601 180 15.4 85.6 510 10 AW520875	177 15.4 85.6 506 14 CB723596	174 15.4 85.6 504 10 BE937951 175 15.4 85.6 504 10 BE833812 176 15.4 05.6 505 14 OBE833812	173 , 15.4 85.6 497 9 AALLI/07	15.4 85.6 495 10 AV815788	15.4 85.6 485 12 BJ061283	15.4 85.6 483 10 AW630245	65 15.4 85.6 478 14 W49749	63 15.4 85.6 478 12 BG961196	161 15.4 85.6 465 10 AW885823 162 15.4 85.6 470 10 BE106945	159 15.4 85.6 452 9 ALO36065 160 15.4 85.6 454 13 BY482752	157 15.4 85.6 439 10 BF117509	155 15.4 85.6 428 14 R58670 156 15.4 85.6 436 9 R1759475	15.2 15.4 85.6 424 14 W30204 15.3 15.4 85.6 427 14 CF112695 15.4 15.4 85.6 428 9 A1909705	151 15.4 85.6 424 12 BI817349

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REFERENCE
AUTHORS
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AA296373/c
                                                        JOURNAL
MEDLINE
PUBMED
COMMENT
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MEDLINE
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                                                                                                                                 Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

El (bases 1 to 23)

El (bases 1 to 23)

Adams,M.D. Kerlavage,A.R., Fleischmann,R.D., Fuldner,R.A.,

Bult,C.J., Lee,N.H., Kirkness,E.F., Weinstock,K.G., Gocayne,J.D.,

White,O., Sutton,G.T.R., Cotton,M.D., Earle-Hughes,J., Fine,L.D.,

Fitzgerald,L.M., Fitzhugh, W.M., Fritchman,J.L., Geoghagen,N.S.,

Glodek,A., Gnehm,C.L., Hanna,M.C., Hedblom,E., Hinkle,P.S.Jr.,

Kelley,J.M., Kelley,J.C., Liu,L.-I., Marmaros,S.M., Merrick,J.M.,

Moreno-Palanques,R.F., McDonald,L.A., Nguyen,D.T., Pelligrino,S.M.,

Fhillips,C.A., Ryder,S.E., Scott,J.L., Saudek,D.M., Shirley,R.,

Small,K.V., Spriggs,T.A., Utterback,T.R., Weidman,J.F., Li,Y.,

Small,K.V., Spriggs,T.A., Utterback,T.R., Weidman,J.F., Li,Y.,

Bednarik,D.P., Cao,L., Cepeda,M.A., Coleman,T.A., Collins,E.J.,

Bednarik,D.P., Greene,J.M., Gruber,J., Hudson,P., Kim,A.K.,

Kozak,D.L., Kunsch,C., Hungjun,J., Li,H., Meissner,P.S., Olsen,H.,

Kozak,D.L., Kunsch,C., Hungjun,J., Li,H., Meissner,P.S., Olsen,H.,

Raymond,L., Wei,Y.F., Wing,J., Xu,C., Yu,G.L., Ruben,S.M.,

Dillion,P.J., Fannon,M.R., Rosen,C.A., Haseltine,M.A., Fields,C.,

Fraser,C.M. and Venter,J.C.

Initial assessment of human gene diversity and expression patterns

based upon 8 million nucleotides of cDNA sequence

NE AG02528
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Contact: Yutaka Suzuki
Contact: Yutaka Suzuki
Department of Virology
Institute of Medical Science, University of Tokyo
4-6-1, Shirokanedai, Minatoku, Tokyo 108-8639, Japan
Email: ysuzuki@ims.u-tokyo.ac.jp
Suzuki,Y., Yoshitomo-Nakagawa,K., Maruyama,K., Suyama,A. and
Sugano,S. Construction and characterization of a full
length-enriched and a 5'-end-enriched cDNA library. Gene 200 (1-2),
149-156 (1997)
                            Other_ESTs: THC168370
Contact: Kerlavage, AR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AA296373 203 bp mRNA linear EST 18-APR-19
EST10934 Umbilical vein endothelial cells II Homo sapiens cDNA 5'
end similar to transformation-associated protein p53, mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This clone was obtained from a 'full length-enriched' cDNA library constructed by 'Oligo-Capping' method. The coding region starts from the 50 bp upstream to the 3'-end.
Location/Qualifiers
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Genomics 64 (3), 286-297 (2000)
20221373
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Bioinformatics
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/mol_type="mRNA"
/db_xref="taxon:9606"
/clone="kaia4737"
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CB117906/c
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Best Local :
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Tel: 3018699056
Fax: 3018699423
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  For clone availability, additional sequence and evinformation related to this EST, please check the Index (http://www.tigr.org/tdb/hgi/hgi.html)
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Genome Research Center
Korea Research Institute of Bioscience & Biotechnology
Schoeun-dong Yuseong-gu, Daejeon 305-333, South Korea
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CB117906 230 bp mRNA linear EST 28-JAN-2003 K-EST0163982 B1T694954 Homo sapiens cDNA clone B1T694954-4-F11 5',
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Seq primer:
                                                                                                                                                                                                                                                                                                                        Email: yongsung@mail.kribb.re.kr
Plate: 4 row: P column: 11
High quality sequence stop: 230.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                        Tel: +82-42-860-4470
Fax: +82-42-860-4409
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          21C Frontier Korean EST Project 2001
Unpublished (2002)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Kim, N.S., Hahn, Y., Oh, J.H., Lee, J.Y., Ahn, H.Y., Chu, M.Y., Kim, M.R., Oh, K.J., Cheong, J.E., Sohn, H.Y., Kim, J.M., Park, H.S., Kim, S. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 230)

Kim, N.S., Hahn, Y., Oh, J.H., Lee, J.Y., Ahn, H.Y., Chu, M.Y., Kim, M.R.
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BST.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Kim YS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens (human)
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/mol type="mxNA"
/db_xref="ATCC (inhost):194290"
/db_xref="teaxon:9606"
/cell type="endothelial cell"
/dev_stage="fetus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /clone lib="Umbilical vein endothelial cells II" /note="Organ: umbilical vein; Vector: pBluescript SK-; Site_1: EcoRI; Site_2: XhoI"
/lab_host="Top10F'"
/clome lib="B17694954"
/clome lib="B17694954"
/clome lib="B17694954"
/note="Organ: Brain; Vector: pCNS-D2; Site 1: EcoRI;
/note="Organ: Brain; Vector: pCNS-D2; Site 1: EcoRI;
Site 2: NotI; The poly (A)+ RNA was dephosphorylated with bacterial alkaline phosphatase (BAP) and then decapped with tabacco acid pyrophosphatase (TAP). The decapped intact mRNA was ligated with DNA-RNA linker including EcoRI site by treatment of T4 RNA ligase and the first
                                                                                                                                                                                                                                    /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
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                                                                                                                                                                                          Sex="M"
                                                                                                                                                                                                            clone="B1T694954-4-F11"
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Pred. No. 4.2e+02;
, Mismatches 0;
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H97230/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Büteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

(bases 1 to 234)

Hillier, L., Clark, N., Dubuque, T., Elliston, K., Hawkins, M.,
Holman, M., Hultman, M., Kucaba, T., Le, M., Lennon, G., Marra, M.,
Parsons, J., Rifkin, L., Rohlfing, T., Soares, M., Tan, F.,
Trevaskis, E., Waterston, R., Williamson, A., Wohldmann, P. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              234 bp mRNA linear E yx04g12.r1 Soares melanocyte 2NbHM Homo sapiens cDNA IMAGE.260806 5' similar to gb:X54156_rnal CELLULAR TU P53 (HUMAN); mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The WashU-Merck EST Project Unpublished (1995) Contact: Wilson RK
                                                                                                                                                                                                                                                                                                                                                                                                        High quality sequence starts: 1
High quality sequence stops: 1
Source: IMAGE Consortium, LLNL
This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (info@image.llnl.gov) for further information.
Trace considered overall poor quality
Insert Length: 2246 Std Error: 0.00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Washington University School of Medicine 4444 Forest Park Parkway, Box 8501, St.
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                                                                                                                                                                                                                                                                                                                                                                                       Seq primer: M13RP1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Email: est@watson.wustl.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Tel: 314 286 1800
Fax: 314 286 1810
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ity sequence stop: 1. Location/Qualifiers
                                                                                                                                                                                                                                               /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="GDB:3870448"
                                                                                                                                                                                                           'db_xref="taxon:9606"
'clone="IMAGE:260806"
                                                                                                                                                                                     sex="Male"
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Pred. No. 4.4e+02;
0; Mismatches 0;
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Matches 18
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401 bp
UI-HF-BM0-adl-a-03-0-UI.r2 NIH N
IMAGE:3061901 5', mRNA sequence
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NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-remail.nih.gov
Tissue Procurement: ATCC
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                                                                                                                                                                                                   18;
                                                                                                                                                                 1 CCCGGAAGGCAGTCTGGC 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cDNA Library Preparation: Life Technologies, Inc. CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can found through the I.M.A.G.E. Consortium/LLNL at: http://inage.llnl.gov plate: LLAM10387 row: e column: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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EST.
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360 bp mRNA linear EST 13-FI 602377775F1 NIH_MGC_92 Homo sapiens cDNA clone IMAGE:4508539
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                                                                                                                                                                                                                                                                                      /note="Organ: tēstis; Vector: pCMV-SPORT6; Site_1: NotI; Site_2: SalI; Cloned unidirectionally: oligo-dT primed. Average insert size 2.5 kb. Library enriched for full-length clones and constructed by Life Technologies. Note: this is a NIH_MGC Library."
                                                                                                                                                                                                                                                                                                                                                                                /clone="IMAGE:4508539"
/cissue type="embryonal carcinoma, cell line"
/lab host="DH10B (phage-resistant)"
/clone_lib="NIH_MGC_92"
                                                                                                                                                                                                                                                                                                                                                                                                                                                     /mol_type="mRNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         organism="Homo sapiens"
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Pred. No. 4.4e+02;
                                                                                                                                                                                                                    Score 18;
Pred. No.
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Eco RI site shown at the beginning of the sequence.

Tissue Procurement: Louis M. Staudt, M.D., Ph.D.

cDNA Library Preparation: M.B. Soares Lab

cDNA Library Arrayed by: M.B. Soares Lab

DNA Sequencing by: M.B. Soares Lab

Clone distribution: MGC clone distribution information

found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (Dases 1 to 434)

Kim,N.S., Hahn,Y., Ch,J.H., Lee,J.Y., Ahn,H.Y., Chu,M.Y., Kim,M.R.
Oh,K.J., Cheong,J.E., Sohn,H.Y., Kim,J.M., Park,H.S., Kim,S. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 401) NIH-MGC http://mgc.nci.nih.gov/. National Institutes of Health, Mammalian Gene Collection (MGC)
Korea Research Institute of Bioscience & Biotechnology 52 Eceun-dong Yuseong-gu, Daejeon 305-333, South Korea Tel: +82-42-860-4409
                                                                                                                                                                                                                                                                                                                                                           K-EST0206507 L15CKK1 Homo sapiens cDNA clone L15CKK1-48-G06
                                                                    Contact: Kim YS
Genome Research Center
                                                                                                       21C Frontier Korean EST Project 2001 Unpublished (2002)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Unpublished (1999)
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/db_xref="taxon:9606"
/clone="IMAGE:3061901"
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                                                                                                                                                        Oh,J.H., Lee,J.Y., Ahn,H.Y., Chu,M.Y., Kim,M.R.,
E., Sohn,H.Y., Kim,J.M., Park,H.S., Kim,S. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
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Pred. No. 5e+02;
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                                                                                                                                                                                            Korea Research Institute of Bioscience & Biotechnology 52 Eoeun-dong Yuseong-gu, Daejeon 305-333, South Korea Tel: +82-42-860-4470 Fax: +82-42-860-4409
                                                                                                                                                                                                                                                                                                                                                                                                           Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 478)

Kim,N.S., Hahn,Y., Oh,J.H., Lee,J.Y., Ahn,H.Y., Chu,M.Y., Kim,M.R.
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Plate: 48 row: G column: 06
High quality sequence stop: 434.
Location/Qualifiers
                                                                                                                                                   Email: yongsung@mail.kribb.re.kr
Plate: 51 row: E column: 01
                                                                                                                                                                                                                                                                                                        Contact: Kim YS
                                                                                                                                                                                                                                                                                                                          21C Frontier Korean EST Project 2001 Unpublished (2002)
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                                                                                                                                                                                                                                                                                                                                                                                      Cheong, J.E., Sohn, H.Y., Kim, J.M., Park, H.S., Kim, S. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /note="Organ: Liver; Vector: pCNS-D2; Site 1: EcoRI; Site_2: NotI; The poly (A)+ RNA was dephosphorylated w bacterial alkaline phosphatase (BAP) and then decapped
/organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
/clone="S11SNU1-51-E01"
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mol_type="mRNA"
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ne_lib="L15CKK1"
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Pred. No.
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Matches
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18; Conserv
                                                                                                                                                                                                                                                                                                                              Email: genomics@hri.co.jp
HRI human cDNA project; 5'- & 3'-end one pass sequencing: Helix
Research Institute; cDNA library construction: Department of
Virology, Institute of Medical Science, University of Tokyo, and
Helix Research Institute.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Helix Research Institute
1532-3 Yana, Kisarazu, Chiba 292-0812, Japan
Tel: 81-438-52-3975
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Unpublished (2000)
Contact: Takao Isogai
Genomics Laboratory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi, Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.

1 (bases 1 to 513)

Ota, T., Nishikawa, T., Suzuki, Y., Ishi, S., Saio, K., Kawai, Y., Ota, T., Nishikawa, T.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              sequence.
AU129655
AU129655.1 GI:10990009
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AU129655 NT2RP2 Homo sapiens cDNA clone NT2RP2005982 5', mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Isogai, T.
HRI human cDNA project
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens (human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Fax: 81-438-52-3986
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Yamamoto,J., Wakamatsu,A., Nakamura,Y., Nagai,T.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CCCGGAAGGCAGTCTGGC 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /clone lib="SISNUI"
//clone lib="SisNuine phosphosphatase dephosphorylated with Subcarial alkaline phosphatase (BAP) and then decapped with tabacco acid pyrophosphatase (TAP). The decapped intact mRNA was ligated with DNA-RNA linker including Sfil site by treatment of 74 RNA ligase and the first strand cDNA was synthesized with Superscript II using Sfil oligo-dT primer. After first strand synthesis, RNA was degraded by NaOH treatment and cDNA was amplified by PCR reaction. The PCR products were digested with Sfil and cloned into DraIII- digested pME18S-FL3 vector. The obtained cDNA vectors were used for transformation of competent cells E. coll Toplof' by electroporation method. The cDNA libraries constructed by this method are full-length enriched cDNA library."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
/clone_lib="NT2RP2"
/note="Vector: pME18SFL3; mRNA from NT2 neuronal precursor
cells after 2-weeks retinoic acid (RA) induction"
                                                              /cell_type="teratocarcinoma"
/cell_line="NT2"
/clone_lib="NT2RP2"
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/cell_line="SNU-1"
                                                                                                                                                     /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
/clone="NT2RP2005982"
                                                                                                                                                                                                                                                                                                     location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               lab_host="Top10F'"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 100.0%; Score 18; DB 12; 100.0%; Pred. No. 5.3e+02; tive 0; Mismatches 0;
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TITLE

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ORIGIN

ORIGIN

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

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REFERENCE
AUTHORS
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BF794506/c
 SOURCE
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VERSION
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CB135305/c
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Best Local S
Matches 18
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Best Local S
Matches 18
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                                                                                                             CB135305
K-EST0187411
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Louis Staudt, M.D., Ph.D.
CDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 (bases 1 to 521)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BF794506 S21 bp mRNA linear EST 12-JAN-2001 602255758F1 NIH_MGC_85 Homo sapiens cDNA clone IMAGE:4339142 5',
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 CCCGGAAGGCAGTCTGGC 18
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Homo sapiens (human)
Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Plate: LLAM9949 row: c column: High quality sequence stop: 519.
Location/Qualifiers
                                                         CB135305.1
                                                                                             sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                found through the I.M.A.G.E. Consortium/LLNL http://image.llnl.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                        Similarity
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                                                                                                                                                                                                                                                              cccecaaecacrcrec 18
                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                          /clone="IMAGE:4339142"
/tissue type="lymphoma, cell line"
/lab_host="DH10B (phage-resistant)"
/clone lib="NIH_MGC 85"
/note="Organ: lymph; Vector: pCMV-SPORT6; Site 1: Not1;
Site 2: Sall; Cloned unidirectionally; oligo-dT primed.
Average insert size 1.867 kb. Library enriched for
full-length clones and constructed by Life Technologies
Note: this is a NIH_MGC Library."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   organism="Homo sapiens"
/mol_type≈"mRNA"
/db_xref="taxon:9606"
                                                           GI:28102030
                                                                                                                 L5HLK1 Homo
                                                                                                                                                                                                                                                                                                                      100.0%; Score 18; DB 10; 100.0%; Pred. No. 5.4e+02;
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                                                                                                                                   537
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                                                                                                               mRNA linear EST 29-JAN-2003 CDNA clone L5HLK1-32-F04 5', mRNA
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AUTHORS
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JOURNAL
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BM833945/c
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AUTHORS
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Best Local Similarity
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(bases 1 to 537)

Kim, N.S., Hahn, Y., Oh, J.H., Lee, J.Y., Ahn, H.Y., Chu, M.Y., Kim, M.R., Oh, K.J., Cheong, J.E., Sohn, H.Y., Kim, J.M., Park, H.S., Kim, S. and Kim, Y.S.

21C Frontier Korean EST Project 2001
                  Unpublished (2002)
Contact: Kim YS
                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (Dases 1 to 538)

Kim,N.S., Hahn,Y., Oh,J.H., Lee,J.Y., Ahn,H.Y., Chu,M.Y., Kim,M.R., Oh,K.Y., Cheong,J.E., Sohn,H.Y., Kim,J.M., Park,H.S., Kim,S. and
                                                                                                                                                                                                            Homo sapiens (human)
                                                                                                                                                                                                                                                                                                                                             K-EST0108801 S11SNU1 Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Email: yongsung@mail.kribb.re.kr
Plate: 32 row: F column: 04
High quality sequence stop: 537.
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Contact: Kim YS
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                                                                                                                                                                                                                                                                                                                      mRNA sequence.
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                                                  Kim,Y.S.
21C Frontier Korean EST Project 2001
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//Colone_lib="Lifilk1"
//Colone_lib="Lifilk1"
//Colone="Organ: Liver; Vector: pCNS-D2; Site 1: EcoRI;
//Colone="Organ: Liver; Vector: pCNS-D2; Site 1: EcoRI;
Site 2: Noti, The poly (A) + RNA was dephosphorylated with
bacterial alkaline phosphatase (BAP) and then decapped
with tabacco acid pyrophosphatase (TAP). The decapped
intect mRNA was ligated with DNA-RNA linker including
EcoRI site by treatment of T4 RNA ligase and the first
strand cDNA was synthesized from oligo dT-selected mRNA by
priming with dT-tailed vector. The dT-tailed vector was
adjusted to have about 60nt. The cDNA vector was
circularized with E. coli DNA ligase after digestion of
EcoRI which site is also included in vector. An RNA strand
converted to a DNA strand by Okayama-Berg method. The
competent cells E. coli Topolof' by electroporation method.
The cDNA libraries constructed by this method are
full-length enriched cDNA library."
  Research Center
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/mol_type="mRNA"
/db_xref="taxon:9606"
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S11SNU1-59-A11 5',
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CB150761/c
LOCUS
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Best Local :
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                                                                                                                                                                                            Contact: Kim YS
                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
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Korea Research Institute of Bioscience & Biotechnology
52 Eccun-dong Yuseong-gu, Dacjeon 305-333, South Korea
Tel: +82-42-860-4470
Fax: +82-42-860-4470
Fax:
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CBISULT 13
CCBISO761/c
CBISO761/c
CBISO761
CCBISO761
CCBISO755 CISNU17 Homo sapiens cDNA clone CISNU17-34-H06 5',
ACCESSION
CCBISO761
VERSION
CCBISO761
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REFERENCE
AUTHORS
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AU127286/c
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                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (Dases 1 to 554)

Ota, T., Nishikawa, T., Suzuki, Y., Ishii, S., Saito, K., Kawai, Y., Yamamoto, J., Wakamatsu, A., Nakamura, Y., Nagai, T., Sugano, S. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      sequence.
AU127286
                                                                                                                                                                                                                                                                                                           Unpublished (2000)
Contact: Takao Isogai
Genomics Laboratory
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AU127286 NT2RP2 Homo sapiens cDNA clone NT2RP2000945 5', mRNA
                                                                                                                        Email: genomics@hri.co.jp
HRI human cDNA project; 5'- & 3'-end one pass sequencing: Helix
Research Institute; cDNA library construction: Department of
Virology, Institute of Medical Science, University of Tokyo, and
Helix Research Institute.
                                                                                                                                                                                                                                                                 Helix Research Institute
1532-3 Yana, Kisarazu, Chiba 292-0812, Japan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /clone_lib="CISNU17"
//clone="Drgan: Cervix; Vector: pCNS-D2; Site 1: EcoRI;
//note="Organ: Cervix; Vector: pCNS-D2; Site 1: EcoRI;
//note="Organ: Cervix; Vector: pCNS-D2; Site 1: EcoRI;
//note="Organ: Cervix; Vector: pcns dephosphorylated with bacterial alkaline phosphatase (BAP) and then decapped with tabacco acid pyrophosphatase (TAP). The decapped intact mRNA was ligated with DNA-RNA linker including EcoRI site by treatment of T4 RNA ligase and the first strand cDNA was synthesized from oligo dT-selected mRNA by priming with dT-tailed vector. The dT-tailed vector was adjusted to have about 60nt. The cDNA vector was circularized with E. coli DNA ligase after digestion of EcoRI which site is also included in vector. An RNA strand converted to a DNA strand by Okayama-Berg method. The obtained cDNA vectors were used for transformation of competent cells E. coli Toploff by electroporation method. The cDNA libraries constructed by this method are full-length enriched cDNA library."
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/mol_type="mRNA"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cell_type="Epithelial"
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/clone="NT2RP2000945"
                                                                                                       Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           tissue_type="Uterine"
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                                         RESULT 16
BM788590/c
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    LOCUS
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Query Match
Best Local
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                                                                                                                                                                                                                   18;
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  BM788590 :
K-EST0067764 S11SNU1 Homo
                                                                                                                                                                         1 CCCGGAAGGCAGTCTGGC 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Email: genomics@hri.co.jp
HRI human cDNA Project, Sugiyama, T.; Wakamatsu, A.; Irie, R.;
Umezawa, A.; Fukuma, M.; Kusakari, S.; Hata, J.; Ishii, S.; Yamamoto, J.;
Isono, Y.; Saito, K.; Nakamura, Y.; Masuho, Y.; Nagai, K.; Isogai, T.
HRI human cDNA project; cDNA library construction & 5'-end one
pass sequencing: Helix Research Institute.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 (bases 1 to 575)
Imabayashi, H., Mori, T., Gojo, S., Kiyono, T., Sugiyama, T., Irie, R., Isagai, T., Hata, J., Tomoya, Y. and Umezawa, A.
Redifferentiation of dedifferentiated chondrocytes and chondrogenesis of human bone marrow stromal cells via chondrosphere formation with expression profiling by large-scale cDNA analysis Exp. Cell Res. 288 (1), 35-50 (2003)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Contact: Takao Isogai
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1532-3 Yana, Kisarazu, Chiba 292-0812, Japan
Tel: 81-438-52-3975
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens (human)
                                                                                                                                                                                                                                           Similarity
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                                                                                                                                  CCCGGAAGGCAGTCTGGC 115
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                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                          /organism="Homo sapiens"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               575 bp mRNA linear I NIESE2 Homo sapiens cDNA clone NIESE2002780
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/cell_line="NT2"
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/clone_lib="NT2RP2"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      location/Qualifiers
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                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                           Score 18;
Pred. No.
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Pred. No. 5
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590 bp mRNA li
sapiens cDNA clone
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    linear EST 05-1
ne S11SNU1-24-G03
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                       EST 05-MAR-2002
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VERSION
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BM849186/c
                                                                                                     VERSION
KEYWORDS
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                                                                                                                                                                                                                                                                                                                                                                  131
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                                                                                                                                                                                                BM849186 590 bp mRNA linear EST UD-MAK-.
K-EST0129212 S13KWS5 Homo sapiens cDNA clone S13KWS5-66-E06 5',
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21C Frontier Korean EST Project 2001
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1 (bases 1 to 590)

Kim,N.S., Hahn,Y., Oh,J.H., Lee,J.Y., Ahn,H.Y., Chu,M.Y., Kim,M.R., Oh,K.J., Cheong,J.E., Sohn,H.Y., Kim,J.M., Park,H.S., Kim,S. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Plate: 24 row: G column: 03
High quality sequence stop: 590.
Location/Qualifiers
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                         mRNA sequence.
BM849186
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Korea Research Institute of Bioscience & Biotechnology 52 Boeun-dong Yuseong-gu, Daejeon 305-333, South Korea
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Unpublished (2002)
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Plate: 24 row: G column: 03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Tel: +82-42-860-4470
Fax: +82-42-860-4409
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens (human)
                                                                          Homo sapiens (human)
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                                                                                                                                                                                                                                                                                                                                                                  CCCGGAAGGCAGTCTGGC 114
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ilarity 100.0%;
Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /cell Type="Numboblast-like"
/cell lipe="SNU-1"
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/clone lib="SIISNU!"
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Site_2: XhoI; The poly (A) + RNA was dephosphorylated with
bacterial alkaline phosphatase (BAP) and then decapped
with tabacco acid pyrophosphatase (TAP). The decapped
intact mRNA was ligated with DNA-RNA linker including Sfil
site by treatment of T4 RNA ligase and the first strand
cDNA was synthesized with Superscript II using Sfil
oligo-dT primer. After first strand synthesis, RNA was
degraded by NaOH treatment and cDNA was amplified by PCR
reaction. The PCR products were digested with Sfil and
cloned into DraIII- digested pME18S-FL3 vector. The
obtained cDNA vectors were used for transformation of
competent cells E. coll Top10F' by electroporation method.
The cDNA libraries constructed by this method are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
/clone="S11SNU1-24-G03"
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                                                                                                                                 GI:19205585
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AUTHORS
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VERSION
KEYWORDS
SOURCE
ORGANISM
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BM849478/c
LOCUS
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                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 592)

Kim.N.S., Hahn,Y., Oh,J.H., Lee,J.Y., Ahn,H.Y., Chu,M.Y., Kim.M.R
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Oh,K.J., Cheong,J.E., Sohn,H.Y., Kim,J.M., Park,H.S., Kim,S. and
Xim,Y.S.
21C Frontier Korean EST Project 2001
                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       mRNA sequence.
BM849478
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Location/Qualifiers
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Plate: 66 row: E column: 06
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                                                            Contact: Kim YS
                                                                                         21C Frontier Korean EST
Unpublished (2002)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BM849478.1 GI:19205877
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                            Genome Research Center
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Conservative
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/lab host="Top10F"
/clome lib="S13KMS5"
/note="Vector: pCNS; Site_1: EcoRI; Site_2: NotI; The poly
/note="Vector: pCNS; Site_1: EcoRI; Site by
/note="Vector: pNA; NotI; The poly
/note="Vector: primating with primating with
/note="Vector: poly vector was adjusted to have about 60nt. The cDNA vector was adjusted to have about 60nt. The cDNA vector was adjusted to have about 60nt. The cDNA vector was adjusted to have about 60nt. The cDNA strand converted to a DNA strand by Okayama-Berg method. The obtained cDNA vectors
/note="Vector: poly vector: poly vector:
Research Institute
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line="KMS-5"
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Pred. No. 5.6e+02;
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of Bioscience & Biotechnology
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RESULT 19
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Best Local Similarity
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Plate: 69 row: A column: 07
High quality sequence stop: 592.
                                                                                                                                                                                                                        Genome Research Center
Korea Research Institute of Bioscience & Biotechnology
52 Eoeun-dong Yuseong-gu, Daejeon 305-333, South Korea
Tel: +82-42-860-4470
Fax: +82-42-860-4409
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 (bases 1 to 602)

Kim,N.S., Hahn,Y., Oh,J.H., Lee,J.Y., Ahn,H.Y., Chu,M.Y., Kim,M.R., Ch,K.J., Cheong,J.E., Sohn,H.Y., Kim,J.M., Park,H.S., Kim,S. and Kim,Y.S.
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                                                                                                                                              Email: yongsung@mail.kribb.re.kr
Plate: 57 row: B column: 08
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Unpublished (2002)
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                                                                 quality sequence stop: 602.
Location/Qualifiers
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//clone lib="School"
//cl
/organism="Homo sapiens"
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RESULT 20
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Best Local
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Xim,N.S., Hahn,Y., Oh,J.H., Lee,J.Y., Ahn,H.Y., Chu,M.Y., Kim,M.R.,

Oh.K.J., Cheong,J.E., Sohn,H.Y., Kim,J.M., Park,H.S., Kim,S. and
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21C Frontier Korean
Unpublished (2002)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                             High quality sequence stop: 606.
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Plate: 23 row: E column: 10
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/clone_lib="Sl1SNU1"
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                                                                            tissue_type="myeloma"
cell_line="KMS-5"
                                                                                                                                                                                                                                     ocation/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    tissue_type="Stomach"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EST Project 2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 18; DB 12;
Pred. No. 5.6e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            dq 909
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 602;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        linear EST 04-MAR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0,
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BM772261/c
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             source
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                132
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (Dases 1 to 606)

Kim,N.S., Hahn,Y., Oh,J.H., Lee,J.Y., Ahn,H.Y., Chu,M.Y., Kim,M.R., Oh,K.J., Cheong,J.E., Sohn,H.Y., Kim,J.M., Park,H.S., Kim,S. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BM772261 606 bp mRNA linear EST 04-MAR-2002 K-EST0056378 S13KMS5s1 Homo sapiens cDNA clone S13KMS5s1-1-B04 5',
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 CCCGGAAGGCAGTCTGGC 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Kim, Y.S.
21C Frontier Korean EST Project 2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BM772261
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Tel: +82-42-860-4470
Fax: +82-42-860-4409
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Korea Research Institute of Bioscience & Biotechnology 52 Eoeun-dong Yuseong-gu, Daejeon 305-333, South Korea
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens (human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BM772261.1 GI:19101876
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Email: yongsung@mail.kribb.re.kr
Plate: 1 row: B column: 04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Genome Research Center
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Contact: Kim YS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CCCGGAAGGCAGTCTGGC 115
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     quality sequence stop: 606 Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
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/clone lib="$13KMS5s1"
/note="Vector: pCNS; Site 1: EcoRI; Site 2: NotI, The poly note="Vector: pCNS; Site 1: EcoRI; Site 2: NotI, The poly A, RNA was dephosphorylaTed with bacterial alkaline phosphatase (BAP) and then decapped with tabacco acid pyrophosphatase (TAP). The decapped intact mRNA was ligated with DNA-RNA linker including EcoR I site by treatment of T4 RNA ligase and the first strand cDNA was synthesized from oligo dT-selected mRNA by priming with dT-tailed vector was adjusted to have about 60nt. The cDNA vector was circularized with E. coli DNA ligase after digestion of EcoRI which site is
                                                                                                                                                                                                                                                                                                                                                                                                     /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
/clone="S13KWS5s1-1-B04"
                                                                                                                                                                                                                                                                                                                       lab_host="Top10F'"
                                                                                                                                                                                                                                                                                                                                                                                   tissue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             . 606
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                                                                                                                                                                                                                                                                                                                                               e_type="myeloma"
line="KMS-5"
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Pred. No. 5.6e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
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SOURCE

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Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ш
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K-EST0191254 L15CKK1 Homo sapiens cDNA clone L15CKK1-29-B07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Email: yongsung@mail.kribb.re.kr
Plate: 29 row: B column: 07
High quality sequence stop: 612.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Korea Research Institute of Bioscience & Biotechnology 52 Eceun-dong Yuseong-gu, Daejeon 305-333, South Korea Tel: +82-42-860-4470 Fax: +82-42-860-4409
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           21C Frontier Korean EST Project 2001 Unpublished (2002)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Kim, N.S.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CB138393.1 GI:28110037
CB138393.1 GI:28110037
EST.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (bases 1 to 612)
m,N.S., Hahn,Y., Oh,J.H., Lee,J.Y., Ahn,H.Y., Chu,M.Y., Kim,M.R.,
K.J., Cheong,J.E., Sohn,H.Y., Kim,J.M., Park,H.S., Kim,S. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
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/cell line="CK-K1"
//ab host="Top10F'"
//lab host="Top10F'"
/clone lib="L15CKK1"
/clone lib="L15CKK1"
/clone lib="L15CKK1"
/clone lib="L15CKK1"
/note="Organ: Liver; Vector: pCNS-D2; Site_1: EcoRI;
/note="Organ: Liver; Vector: pCNS-D2; Site_1: E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             clone="L15CKK1-29-B07"
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Pred. No. 5.6e+02;
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RESULT 22 CB138393/c LOCUS

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RESULT 24
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VERSION
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BE613913/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOURCE
ORGANISM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      KEYWORDS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DEFINITION
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JOURNAL
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                                                                                                           96
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (Dases 1 to 622)

NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC) Unpublished (1999)

Contact: Robert Strausberg, Ph.D.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CDNA Library Preparation: Life Technologies, Inc.
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
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BE613913.1 GI:9895510
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601504033F1 NIH_MGC_71 Homo:
    AU131823
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                http://image.llnl.gov
Plate: LLAM9713 row: b column: 10
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                                                                                                                                              CCCGGAAGGCAGTCTGGC 18
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                                                                                                        CCCGGAAGGCAGTCTGGC 79
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          quality sequence stop: 610.
Location/Qualifiers
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                                                                                                                                                                                                                                                                                           /tissue_type="leiomyosarcoma"
/lab host="DH10B (phage-resistant)"
/clome_lib="NHH MGC_71"
/clome_Torgan: uterus; Vector: pCMV-SPORT6; Site_1: NotI;
Site_2: SalI; Cloned unidirectionally. Primer: Oligo dT
Average insert size 2.1 kb. "
                                                                                                                                                                                                                                                                                                                                                                                                                                         /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                          clone="IMAGE:3905577"
                                                                                                                                                                                                          100.0%;
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                                                                                                                                                                                                        Score 18; DB 10;
Pred. No. 5.6e+02;
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    TURNA
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    linear
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CB134750/c
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Matches 18
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                                                                                                                  Kim, Y.S.
21C Frontier Korean E
Unpublished (2002)
Contact: Kim YS
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AU131823
AU131823.1
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                     52 Eoeun-dong Yuseong-gu,
Tel: +82-42-860-4470
Fax: +82-42-860-4409
                                         Genome Research Center Genome Research Institute of Bioscience & Biotechnology Korea Research Institute of Bioscience & Biotechnology Foreun-dong Yuseong-gu, Daejeon 305-333, South Korea
                                                                                                                                                                                                Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 649)

Kim,N.S., Hahn,Y., Oh,J.H., Lee,J.Y., Ahn,H.Y., Chu,M.Y., Kim,M.R., Kim,N.S., Hahn,Y., Sohn,H.Y., Kim,J.M., Park,H.S., Kim,S. and Y., Y., Cheong,J.E., Sohn,H.Y., Kim,J.M., Park,H.S., Kim,S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 CCCGGAAGGCAGTCTGGC 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Email: genomics@hri.co.jp
HRI human cDNA project; 5'- & 3'-end one pass sequencing: Helix
Research Institute; cDNA library construction: Department of
Virology, Institute of Medical Science, University of Tokyo, and
Helix Research Institute.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1. (bases 1 to 633)
                                                                                                                                                                                                                                                                                                            Homo sapiens
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CB134750.1
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Fax: 81-438-52-3986
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Helix Research Institute
1532-3 Yana, Kisarazu, Chiba 292-0812, Japan
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Yamamoto,J., Wakamatsu,A., Nakamura,Y.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AU131823 NT2RP3 Homo sapiens cDNA clone NT2RP3003309 5', mRNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Contact: Takao Isogai
Email: yongsung@mail.kribb.re.kr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
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/note="Vector: pME188FL3; mRNA from NT2 neuronal precursor cells after 2-weeks retinoic acid (RA) induction"
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/cell_line="NT2"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /mol_type="mRNA"
/db_xref="taxon:9606"
/clone="NT2RP3003309"
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ura,Y., Nagai,T., Su
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CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
Plate: LLAM10428 row: k column: 12
High quality sequence stop: 652.
Location/Qualifiers
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 657)
11 (bases 1 to 657)
12 (bases 1 to 657)
13 (bases 1 to 657)
14 (bases 1 to 657)
15 (bases 1 to 657)
16 (bases 1 to 657)
17 (bases 1 to 657)
18 (bases 1 to 657)
19 (base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
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/note="Organ: Liver; Vector: pCNS-D2; Site 2: Color place with bacco acid pyrophosphatase (BAP) and then decapped with tabacco acid pyrophosphatase (TAP). The decapped intact mRNA was ligated with DNA-RNA linker including EcoR1 site by treatment of T4 RNA ligase and the first strand cDNA was synthesized from oligo dT-selected mRNA by priming with dT-tailed vector. The dT-tailed vector was adjusted to have about 60nt. The cDNA vector was circularized with E. coli DNA ligase after digestion of EcoR1 which site is also included in vector. An RNA strand converted to a DNA strand by Okayama-Berg method. The obtained cDNA vectors were used for transformation of competent cells E. coli Toploff by electroporation method. The cDNA libraries constructed by this method are full-length enriched cDNA library."
                                                                                                                                          /organism="Homo sapiens"
/mol_type="mRNA"
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/lab_host="Top10F'"
/clone_lib="L5HLK1"
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mol_type≈"mRNA"
(db_xref="taxon:9606"
(clone="L5HLK1-31-F08")
tissue_type="embryonal carcinoma, cell line"
                                               one="IMAGE: 4524419"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     sequence stop: ation/Qualifiers
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Pred. No. 5.7e+02;
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105 CCCGGAAGGCAGTCTGGC
                                                                                                    18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Life Technologies, Inc.
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
plate: LLAM11556 row: a column: 10
                                                1 CCCGGAAGGCAGTCTGGC 18
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National Institutes of Health, Mammalian Gene Collection (MGC)
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1 (bases 1 to 682)
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Location/Qualifiers
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                                                                                                                            Similarity
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                                                                                                                                                                                                                      /clone="IMAGE:5221137"
//lab host="DHIOB"
//lab host="DHIOB"
//clone lib="NHI MGC 120"
//clone lib="NHI MGC 120"
//note="Organ: pooled pancreas and spleen; Vector:
//note="Organ: pooled pancreas and spleen; Vector:
//note="Organ: pooled pancreas and spleen; Vector:
//note="Organ: pooled pancreas and spleen 28 yource anonymous pool of spleen and pancreas from 28 yource anonymous pool of spleen and pancreas from 28 yource anonymous pool of spleen and directionally cloned (EcoRV site is destroyed upon cloning). Average insert size 1.5 kb, insert size range 1-2.5 kb. Library is normalized and enriched for full-length clones and was constructed by C. Gruber (Invitrogen). Research Genetics tracking code 025. Note: this is a NIH_MGC Library."
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/clone lib="NIH MGC 92"
/note="Organ: testis; Vector: pCMV-SPORT6; Site 1: Not1;
/note="Organ: testis; Vector: pCMV-SPORT6; Site 1: Not1;
Site_2: Sall; Cloned unidirectionally; Oligo-dT primed.
Average insert size 2.5 kb. Library enriched for
full-length clones and constructed by Life Technologies.
Note: this is a NIH_MGC Library."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
                                                                                                                            100.0%;
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100.0%; Pred. No. 5.7e+02;
tive 0; Mismatches 0;
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Pred. No. 5.8e+02;
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sapiens cDNA clone IMAGE:5221137 5',
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HRI human cDNA project; 5'- & 3'-end one pass sequencing: Helix
Research Institute; cDNA library construction: Department of
Virology, Institute of Medical Science, University of Tokyo, and
Helix Research Institute.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Helix Research Institute
1532-3 Yana, Kisarazu, Chiba 292-0812, Japan
Tel: 81-438-52-3975
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Genomics Laboratory
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Ota, T., Nishikawa, T., Suzuki, Y., Ishii, S., Saito, K., Kawai, Y., Yamamoto, J., Wakamatsu, A., Nakamura, Y., Nagai, T., Sugano, S. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AU121050 688
AU121050 HEMBB1 Homo sapiens
                                               Saito, K., Yamamoto, J., Nakamura, Suzuki, Y., Sugano, S., Isogai, T.) Cupublished (2000)
                                                                                  HRI human cDNA project (Ota,T., Wakamatsu,A., Ozawa,M., Ishii,S., Saito,K., Yamamoto,J., Nakamura,Y., Nishikawa,T., Nagai,T.,
                                                                                                                                          Ota,T., Wakamatsu,A., Ozawa,M., Ishii,S.,
Nakamura,Y., Nishikawa,T., Nagai,T., Suzu
                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 694)
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Unpublished (2000)
Helix Research Institute
                                 Contact: Takao Isogai
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               Genomics Laboratory
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /organism="Homo sapiens"
/mol type="mRNA"
/db xref="texon:9606"
/clone="HEMBB1001969"
/tissue_type="whole embryo, mai/dev_stage="embryo, 10 weeks"
/clone_lib="HEMBB1"
/note="Vector: pME185FL3"
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CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
Plate: LLCM1958 row: j column: 22
High quality sequence stop: 696.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 (bases 1 to 707)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
Unpublished (1999)
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BM011389
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Email: genomics@hri.co.jp
HRI human cDNA project; 5'- & 3'-end one pass sequencing: Helix
Research Institute; cDNA library construction: Department of
Virology, Institute of Medical Science, University of Tokyo, and
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Fax: 81-438-52-3986
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Mammalia; Eutheria; Primates;
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llarity 100.0%;
Conservative 0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /organism="Homo sapiens"
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/clone="NT2RM4000122"
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/cell_line="NT2"
/tissue_type="neuroblastoma, cell line"
/lab_host="DH10B (phage resistant)"
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/clone lib="NNH MGC 47"
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/note="Organ: brain; Vector: pOTB7; Site_1: XhoI, Site_2:
ECORI, CDNA made by oligo-dT priming. Directionally
cloned into ECORI/XhoI sites using the following 5;
adaptor: GGCACGAG(G). Size-selected >500bp for average
insert size 1.8kb. Library constructed by Ling Hong in
the laboratory of Gerald M. Rubin (University of
California, Berkeley) using ZAP-cDNA synthesis kit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /clone lib="NT2RM4" /note="Vector: pME18SFL3; mRNA from uninduced NT2 neuronal precursor cells"
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/clone="IMAGE:5457909"
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Pred. No. 5.8e+02;
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AU141723/c
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Best Local S
Matches 18
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JOURNAL
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                ORGANISM
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AU131952 NT:
sequence.
AU131952
AU131952.1
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HRI humar
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Unpublished (2000)
Contact: Takao Isogai
Genomics Laboratory
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[ (bases 1 to 723)
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AU141723 THYRO1
                                                                                                                                                                                                                                                                                                                                                                                                                                Email: genomics@hri.co.jp
HRI human cDNA project; 5'- & 3'-end one pass sequencing: Helix
Research Institute; cDNA library construction: Department of
Virology, Institute of Medical Science, University of Tokyo, and
Helix Research Institute:
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1532-3 Yana, Kisarazu, Chiba
Tel: 81-438-52-3975
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          sequence.
AU141723
 Eukaryota; Metazoa; Chordata; Craniata;
                Homo sapiens
                             Homo sapiens (human)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
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                                                                                                                                                                                                                                                                                                       /organism="Homo sapiens"
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/tissue_type="thyroid gland"
/clone_lib="THYRO1"
/note="Vector: pME18SFL3"
                                                                                                NT2RP3
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                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
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                                                         GI:10992306
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                                                                                         724
Homo sapiens
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Pred. No. 5.8e+02;
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clone NT2RP3003525
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Vertebrata;
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  Euteleostomi;
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55', mRNA
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Tel: 81-438-52-3975
Fax: 81-438-52-3986
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                                                                                                                                                                                                                                                                                                                                     Homo sapiens (human)
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llarity 100.0%;
Conservative 0
 /organism≃"Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
/clone="IMAGE:5459560"
                                                                                                                                                                                                                                                                                                                                                                GI:16526756
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cDNA Library Preparation: Ling Hong/Rubin Laboratory cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov plate: LLCM1962 row: o column: 17 High quality sequence stop: 542.
                                                                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 726)

NIH-MGC http://mgc.nci.nih.gov/.

National Institutes of Health, Mammalian Gene Collection (MGC)

Unpublished (1999)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Email: genomics@hri.co.jp
HRI human cDNA project; 5'- & 3'-end one pass sequencing: Helix
Research Institute; cDNA library construction: Department of
Virology, Institute of Medical Science, University of Tokyo, and
Helix Research Institute.
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Helix Research Institute
Helix Research Institute
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(bases 1 to 724)
Ota, T., Nishikawa, T., Suzuki, Y., Ishii, S., Saito, K., Kawai, Y.,
Yamamoto, J., Wakamatsu, A., Nakamura, Y., Nagai, T., Sugano, S. and
                                                                                                                                                                                                                                                          Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
                                                                                                                                                                                                                                                                                                                   Contact: Robert Strausberg, Ph.D.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /clone_lib="NT2RP3"
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cells after 2-weeks retinoic acid (RA) induction"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /organism="Homo sapiens"
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/cell_line="NT2"
Location/Qualifiers
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Pred. No.
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SOURCE
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AU141300/c
RESULT 35
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JOURNAL
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Best Local (
                                                                                                                                         Matches
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AU141300
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AU141300.1
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Genomics Laboratory
Helix Research Institute
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1 (bases 1 to 729)

Ota,T., Nishikawa,T., Suzuki,Y., Ishii,S., Saito,K., Kawai,Y.,
Yamamoto,J., Wakamatsu,A., Nakamura,Y., Nagai,T., Sugano,S. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               18;
                                                                                                                                         18;
                                                                                                                                                                                                                                                                                                                                                                                                 Email: genomics@hri.co.jp

HRI human cDNA project; 5'- & 3'-end one pass sequencing: Helix
Research Institute; cDNA library construction: Department of
Virology, Institute of Medical Science, University of Tokyo, and
Helix Research Institute.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    HRI human cDNA project 
Unpublished (2000)
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                                                                                                                                                          Similarity
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                                                                                        CCCGGAAGGCAGTCTGGC 18
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                                                                                                                                       Conservative
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/clone_lib="NIH_MGC_47"
/clone_lib="NIH_MGC_47"
/clone_lib="NIH_MGC_47"
/cloned into EcoRI/KhoI sites using the following 5'
adaptor: GCCACCAG(G). Size-selected >500bp for average
insert size 1.8kb. Library constructed by Ling Hong in
the laboratory of Gerald M. Rubin (University of
California, Berkeley) using ZAP-cDNA synthesis kit
(Stratagene) and Superscript II RT (Life Technologies).
Note: this is a NIH_MGC Library."
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THYRO1 Homo sapiens
                                                                                                                                                                                                                                                                                                                     /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                      /note="Vector: pME18SFL3"
                                                                                                                                                                                                                                                           /clone="THYRO1000362"
/tissue_type="thyroid gland"
/clone_lib="THYRO1"
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                                                                                                                                   100.0%; Score 18; DB 9; I
100.0%; Pred. No. 5.9e+02;
tive 0; Mismatches 0;
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                                                                                                                                                                            DB 9; Length 729;
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BE273022/c
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Best Local Similarity
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 744)

NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC) Unpublished (1999)

Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: DCTD/DTP
                                                                                                                                                                                                                                                BE273022 744 bp mRNA linear EST 13-JUL-2000 601171343F1 NIH MGC_14 Homo sapiens cDNA clone IMAGE:3544714 5',
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High quality sequence stop: 715.
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1 (bases 1 to 732)
1 (bases 1 to 732)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
Unpublished (1999)
                                                                                                                                                   Homo sapiens
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BE273022
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Tissue Procurement: DCTD/DTP
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ilarity 100.0%;
Conservative 0
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/mol_type="mRNA"
/db_xref="taxon:9606"
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BM048815/c
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ORGANISM
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                                                                                                                                                                                                                                                                                       http://image.llnl.gov
Plate: LLCM1955 row: h column: 16
High quality sequence stop: 715.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov Tissue Procurement: DCTD/DTP
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 750)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
Unpublished (1999)
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BM048815
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                                                                                                                                                                                                                                                                                                                                                           DNA Sequencing by: Incyte Genomics, Inc.

Clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at:
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/tissue_type="renal cell adenocarcinoma"
/tissue_type="renal cell adenocarcinoma"
/lab host="DHIOB (phage-resistant)"
/clone=lib="NIH MGC 14"
/clone=lib="NIH MGC 14"
/clone="Crgan: kidney; Vector: pOTB7; Site_1: XhoI; Site_2:
ECORI; CDNA made by oligo-dT priming. Directionally
cloned into ECORI/XhoI sites using the following 5'
adaptor: GGCACGAG(G). Size-selected >500bp for average
insert size 1.8kb. Library constructed by Ling Hong in
the laboratory of Gerald M. Rubin (University of
California, Berkeley) using ZAP-CDNA synthesis kit
(Stratagene) and Superscript II RT (Life Technologies)."
/tissue_type="carcinoma, cell line"
/lab_host="pH10B (phage-resistant)"
/clone_lib="WIH_MGC_40"
/clone_lib="WIH_MGC_40"
/note="Organ: prostate; Vector: pOTB7; Site_1: XhoI;
Site_2: EcoRI; cDNA made by oligo-dT priming.
                                                                                                                                                         organism="Homo sapiens"
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/db_xref="taxon:9606"
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/mol_type="mRNA"
/db_xref="taxon:9606"
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Pred. No. 5.9
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. 5.9e+02;
ches 0;
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one IMAGE:5456703 5'
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                                       1 CCCGGAAGGCAGTCTGGC 18
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1 (Dases 1 to 758)

NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
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CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
Plate: LLAM11558 row: p column: 11
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Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Life Technologies,
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BI839050.1 GI:15950600
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                                                                                                                                                                                /clone lib="NIH MGC 120"
/clone lib="NIH MGC 120"
/note="Organ: pooled pancreas and spleen; Vector:
/note="Organ: pooled pancreas and spleen; Vector:
pCMV-SPORT6; Site 1: Not1; Site 2: EcoRV (destroyed); RNA
source anonymous pool of spleen and pancreas from 28 yo
male. Library is oligo-dT primed and directionally cloned
(EcoRV site is destroyed upon cloning). Average insert
size 1.5 kb, insert size range 1-2.5 kb. Library is
normalized and enriched for full-length clones and was
constructed by C. Gruber (Invitrogen). Research Genetics
tracking code 025. Note: this is a NIH_MGC Library."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Directionally cloned into EcoRI/XhoI sites using the following 5' adaptor: GGCAGGAG(G). Library constructed by Ling Hong in the laboratory of Gen. Rubin (University of California, Berkeley) using ZAP-cDNA synthesis kit (Stratagene) and Superscript II RT (Life Technologies). Note: this is a NIH_MGC Library."
                                                                                                                                                                                                                                                                                                                                                                                                                               /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
/clone="IMAGE:5222266"
                                                                                                                                                                                                                                                                                                                                                                                                             ab_host="DH10B"
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                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 pro 765)

NIH-MGC http://mgc.nci.nih.gov/.

National Institutes of Health, Mammalian Gene Collection (MGC)

Unpublished (1999)

Contact: Robert Strausberg, Ph.D.
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EST.
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BP 191 91006 EVRY cedex - France
BR 191 91006 EVRY cedex - France
Email: seqref@genoscope.cns.fr, Web: www.genoscope.cns.fr
Library was constructed by Life Technologies, a division of
Invitrogen. This sequence belongs to sequence cluster 8550.
more information about this cluster, see
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603622210F1 NIH_MGC_40 Homo (
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cgi-bin/cluster.cgi?seq=CSODH003AH02QP1&cluster=8550.f. Contact
Feng Liang Email : fliang@lifetech.com URL :
http://fulllength.invitrogen.com/ InVitroden Corporation 1600
Faraday Avenue Genoscope sequence ID : CSODH003AH02QP1.
Location/Qualifiers
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                                                                                  Email:
                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              mRNA sequence.
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    CDNA Library Preparation:
                                     mail: cgapbs-r@mail.nih.gov
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/tissue type="T CELLS (JURKAT CELL LINE)"
/cell_line="UJURKAT CELL LINE"
/clone_lib="Homo sapiens T CELLS (JURKAT CELL LINE)"
/note="Vector: pCMVSPORT 6; 1st strand cDNA was primed with a NotI-oligo(dT) primer. Five prime end enriched, double-strand cDNA was digested with Not I and cloned into the Not I and ECCRV sites of the pCMVSPORT 6 vector.
Library was not normalized."
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/mol_type="mRNA"
/db_xref="taxon:9606"
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Ling Hong/Rubin Laboratory
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cDNA clone IMAGE:5447739 5',
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BM046401/c
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Matches 18
                                                                                                                                                                                                                      source
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1 (Dases 1 to 765)

NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
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                                                                                                                                                                                                                                                                 Plate: LLCM1945 row: d column: High quality sequence stop: 544.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Email: cgapbs-r@mail.nih.gov
Tissue Procurement: DCTD/DTP
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                                                                                                                                                                                                                                                                                                              found through the I.M.A.G.E. Consortium/LLNL http://image.llnl.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                              CDNA Library Preparation: Ling Hong/Rubin Laboratory CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can be
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(larity 100.0%;
Conservative 0,
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/clone lib="NIH MGC 40"
/clone lib="NIH MGC 40"
/note="Organ: prostate; Vector: pOTB7; Site_1: XhoI;
Site_2: EcoRI; cDNA made by oligo-dT priming.
Directionally cloned into EcoRI/XhoI sites using the following 5' adaptor: GGACGAGG(). Library constructed by Ling Hong in the laboratory of Gerald M. Rubin (University of California, Berkeley) using ZAP-cDNA synthesis kit (Stratagene) and Superscript II RT (Life Technologies).
Note: this is a NIH_MGC Library."
/tissue_type="carcinoma, cell line"
/lab host="DHIOB (phage resistant)"
/clone_lib="WIH_MGC_40"
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/note="Organ: prostate; Vector: pOTB7; Site_1: XhoI;
Site_2: EcoRI; cDNA made by oligo-dT priming.
                                                                                                                                               organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                             ocation/Qualifiers
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mol_type="mRNA"
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                                                                                                                           clone="IMAGE:5452758"
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Pred. No.
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JOURNAL COMMENT
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AUTHORS
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ORGANISM
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BM048326/c
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VERSION
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RESULT 43
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                                                                                                                                                                         Query Match
Best Local (
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                                                               115 CCCGGAAGGCAGTCTGGC 98
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (Dases 1 to 765)

NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
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603625546F1 NIH_MGC_40 Homo sapiens
                                                                                                     1 CCCGGAAGGCAGTCTGGC 18
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High quality sequence stop: 341.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CDNA Library Preparation: Ling Hong/Rubin Laboratory CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov
Tissue Procurement: DCTD/DTP
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BM048326.1 GI:16777593
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Unpublished (1999)
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                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                         /clone="IMAGE:5452187"
/tissue_type="carcinoma, cell line"
/tissue_type="carcinoma, cell line"
/lab_host="DHIOB (phage-resistant)"
/clone_lib="NIH_MGC_40"
/clone_lib="NIH_MGC_40"
/clone_lib="NIH_MGC_40"
/clone_lib="Corror priming.
Site_2: EcoRI; cDNA made by ollood friming.
Directionally_cloned into EcoRI/XhoI sites using the following 5' adaptor: GGCACGAG(G). Library constructed by Ling Hong in the laboratory of Gerald M. Rubin (University of California, Berkeley) using ZAP-cDNA synthesis kit (Stratagene) and Superscript II RT (Life Technologies).
Note: this is a NIH_MGC Library."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
                                                                                                                                                 100.0%; Score 18; DB 12
100.0%; Pred. No. 6e+02;
tive 0; Mismatches
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cone IMAGE:5452187
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REFERENCE
AUTHORS
TITLE
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COMMENT
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AUTHORS
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ORGANISM
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ORGANISM
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DEFINITION
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AU120569/c
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KEYWORDS
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Best Local S
Matches 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       110 CCCGGAAGGCAGTCTGGC 93
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ch 100.0%;
l Similarity 100.0%;
18; Conservative 0;
HRI human cDNA project
Unpublished (2000)
Contact: Takao Isogai
                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 781)
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                                                                                                                                                                                                                                                                            AU120569
AU120569.1 GI:10935804
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AU120569 HEMBB1 Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CDNA Library Preparation: Ling Hong/Rubin Laboratory CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: http://inage.llnl.gov Plate: LLCM1437 row: 1 column: 03
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1 (bases 1 to 767)

NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
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EST.
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BG481823
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602529214F1 NIH_MGC_21 Homo sapiens
                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                              Homo sapiens (human)
                                                                                                                                                                                                                                                                                                                        sequence.
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Location/Qualifiers
1. .767
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Tissue Procurement: ATCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens (human)
                                                                                                                Ota, T., Nishikawa, T., Suzuki, Y.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Unpublished (1999)
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                                                                                     Yamamoto, J., Wakamatsu, A., Nakamura, Y.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CCCGGAAGGCAGTCTGGC 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /clone="IMAGE:4652690"
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/lab_host="DHIOB (phage-resistant)"
/lab_host="DHIOB (phage-resistant)"
/clone lib="NIH MGC 21"
/clone lib="NIH MGC 21"
/clone lib="NIH MGC 21"
/note="organ: placenta; Vector: pOTB7; Site_1: XhoI;
/site_2: EcoRI; cDNA made by oligo-dT priming.
Sitecinnally cloned into EcoRI(XhoI sites using the prectionally cloned into EcoRI(XhoI sites using the following 5 adaptor: GGCACGAG(G). Size-selected >500bp for average insert size 1.8kb. Library constructed by Ling Hong in the laboratory of Gerald M. Rubin (University of California, Berkeley) using ZAP-cDNA synthesis kit (Stratagene) and Superscript II RT (Life Technologies)."
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/mol_type="mRNA"
/db_xref="taxon:9606"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 18; DB 12
Pred. No. 6e+02;
; Mismatches
                                                                                                                                                                                                                                                                                                                                         bp mRNA linear EST 01-AUG-2002 cDNA clone HEMBB1000956 5', mRNA
                                                                                                             Ishii, S., Saito, K., Kawai, Y.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 12;
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s cDNA clone IMAGE:4652690 5',
                                                                                       Nagai,T.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 767;
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                                                                                       Sugano, S. and
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FEATURES

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Query Match
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HRI human cDNA project; 5'- & 3'-end one pass sequencing: Helix
Research Institute; cDNA library construction: Department of
Virology, Institute of Medical Science, University of Tokyo, and
Helix Research Institute.
                                                                                                                                                                                                                                                                                                                                                                  CDNA Library Preparation: Life Technologies, Inc.
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
http://image.llnl.gov
plate: LLAM11438 row: b column: 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Life Technologies.
cDNA Library Preparation: Life Techno.
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1 (Dases 1 to 808)

NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
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B1821174
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Tel: 81-438-52-3975
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Helix Research
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/lab_host="DH10B"
/clone lib="NNIH MGC 115"
/clone lib="NNIH MGC 115"
/note="Organ: pooled brain, lung, testis; Vector:
/note="Organ: pooled brain, lung, testis; Vector:
pCMV-SPORT6; Site 1: NotI; Site 2: EcoRV (destroyed); RNA
source anonymous pool of 6 male brains, age range 23-27; 1
male lung, age 27; and 1 male testis, age 69. Library is
oligo-dT primed and directionally cloned (EcoRV site is
oligo-dToprimed and directionally cloned (EcoRV site is
destroyed upon cloning). Average insert size 1.8 kb,
insert size range 1-3 kb. Library is normalized and
                                                                                                                                                                                                         /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
/clone="IMAGE:5175858"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       tissue type="whole embryo, m'dev_stage="embryo, 10 weeks"
clone_lib="HEMBB1"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            organism="Homo sapiens"
(mol_type="mRNA"
(db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           clone="HEMBB1000956"
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sarazu, Chiba 292-0812, Japan
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                        RESULT 47
BU902747/c
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COMMENT
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BI905893/c
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RESULT 45 BI821174/c

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Matches

REFERENCE AUTHORS

TITLE

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SOURCE ORGANISM KEYWORDS

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1 CCCGGAAGGCAGTCTGGC 18
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Plate: LLMN11532 row: h column: 07
High quality sequence stop: 778.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Life Technologies, Inc.
CDNA Library Preparation: Life Technologies, Inc.
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 815)

NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 CCCGGAAGGCAGTCTGGC 18
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18; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Unpublished (1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapıens (human)
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                                                                                                                                                                                                                                    /clone_lib="NIH_MGC_118"
/clone_TVector: pCMV-SPORT6; Site 1: NotI; Site 2: EcoRV
/note="Vector: pCMV-SPORT6; Site 1: NotI; Site 2: EcoRV
(destroyed); RNA source leukocytes from anonymous pool of
non-activated adult donors. Library is oligo-dT primed
and directionally cloned (EcoRV site is destroyed upon
cloning). Average insert size 1.7 kb, insert size range
1.2-3 kb. Library is normalized and enriched for
full-length clones and was constructed by C. Gruber
(Invitrogen). Research Genetics tracking code 027. Note:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gruber (Invitrogen). Research Genetics tracking 021. Note: this is a NIH_MGC Library."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /mol_type="mRNA"
/db_xref="taxon:9606"
/clone="IMAGE:5212086"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 tissue_type="leukocyte"
'lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   organism="Homo sapiens"
                                                                                          100.0%; Score 18; DB 12; 100.0%; Pred. No. 6.1e+02; O. Mismatches 0;
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Perfect score:
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                                                                                                                                                                                                                                                                         Score
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seq length:
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                                                                                                                                                                                                                                                                                                                                                                                                               Published Applications NA:*

1: /cgn2_6/ptodata/2/pubpna/US07_PUBCOMB.seq:*

2: /cgn2_6/ptodata/2/pubpna/US06_NEW_PUB.seq:*

3: /cgn2_6/ptodata/2/pubpna/US06_NEW_PUB.seq:*

4: /cgn2_6/ptodata/2/pubpna/US06_PUBCOMB.seq:*

5: /cgn2_6/ptodata/2/pubpna/US08_PUBCOMB.seq:*

6: /cgn2_6/ptodata/2/pubpna/US08_NEW_PUB.seq:*

7: /cgn2_6/ptodata/2/pubpna/US08_NEW_PUB.seq:*

8: /cgn2_6/ptodata/2/pubpna/US08_PUBCOMB.seq:*

9: /cgn2_6/ptodata/2/pubpna/US09A_PUBCOMB.seq:*

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11: /cgn2_6/ptodata/2/pubpna/US09A_PUBCOMB.seq:*

12: /cgn2_6/ptodata/2/pubpna/US09A_PUBCOMB.seq:*

13: /cgn2_6/ptodata/2/pubpna/US09A_PUBCOMB.seq:*

14: /cgn2_6/ptodata/2/pubpna/US09A_PUBCOMB.seq:*

15: /cgn2_6/ptodata/2/pubpna/US09_PUBCOMB.seq:*

16: /cgn2_6/ptodata/2/pubpna/US10A_PUBCOMB.seq:*

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17: /cgn2_6/ptodata/2/pubpna/US09_PUBCOMB.seq:*
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Listing first 200 summaries
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US-09-776-655-22
US-09-7732-384-8
US-09-732-384-8
US-09-732-384-8
US-09-829-922-1
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US-10-160-290-1
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US-10-160-290-1
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US-10-160-290-1
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US-10-160-29-121-4
8
US-10-160-29-131-4
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US-10-160-29-131-4
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US-10-430-503-16
US-10-077-176-59
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20383, A
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Publication No. US20030166588A1

Publication No. US20030166588A1

APPLICANT: IVERSEN, Patrick L.

APPLICANT: Hudziak, Robert

TITLE OF INVENTION: Splice-Region Antisense Composition and TITLE OF INVENTION: Method

FILE REFERENCE: 0450-0037.30

CURRENT APPLICATION NUMBER: US/09/848,868

CURRENT FILING DATE: 2001-05-04

PRIOR APPLICATION NUMBER: US 60/202,376

PRIOR A
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: TYPE: DNA
: ORGANISM: Artificial Sequence
: FEATURE:
: OTHER INFORMATION: antisense
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Sequence 2883, Ap
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Sequence 3, Appli
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US-09-776-695-22/c
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; Patent No. US2002068283A1
; GENERAL INFORMATION:
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US-09-848-868-36
) Sequence 36, Application US/09848868
; Publication No. US20030166588A1
; GENERAL INFORMATION:
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/848,868
CURRENT FILING DATE: 2001-05-04
PRIOR APPLICATION NUMBER: US 60/202,376
PRIOR FILING DATE: 2000-05-04
NUMBER OF SEQ ID NOS: 54
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Hudziak, Robert
TITLE OF INVENTION: Splice-Region Antisense Composition and
TITLE OF INVENTION: Method
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Iversen, Patrick L. APPLICANT: Hudziak, Robert
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ORGANISM: Artificial Sequence
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MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FRATSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/776,695
FILING DATE: 26-Feb-2001
CLASSIFICATION: «Unknown»
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
ADDRESSEE: Banner & Witcoff
STREET: 1001 G Street, NW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: NUCLEIC ACIDS ENCODING P53
MUTATIONS WHICH SUPPRESS P53 CANCER MUTA- TIONS
                 TELECOMMUNICATION INFORMATION: TELEPHONE: 202-508-9100
                                                                                                           APPLICATION NUMBER: 08/795,006
FILING DATE: 2001-03-28
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Boeke, Jef
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          18;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CCCGGAAGGCAGTCTGGC 18
                                                 NAME: Kagan, Sarah A
REGISTRATION NUMBER: 32141
REFERENCE/DOCKET NUMBER: 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CCCGGAAGGCAGTCTGGC 18
                                                                                                                                                                                                                                                                                                                                                                                                                                         TY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%; Score 18; DB 10; Length 36; Llarity 100.0%; Pred. No. 12; Conservative 0; Mismatches 0; Indels
202-508-9299
                                                                                                                                                                                                                                                                                                                                                                                                       USA
                                                       01107.03170
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LENGTH: 1303
; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-397-635-6
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US-09-732-384-8/c
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-732-384-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Loc
Matches
                                                                       CURRINT APPLICATION NUMBER: US/09/732,384
CURRENT FILING DATE: 2000-12-07
PRIOR APPLICATION NUMBER: 60/169,816
PRIOR FILING DATE: 1999-12-08
NUMBER OF SEQ ID NOS: 10
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 8
SEQ ID NO 8
                                                                                                                                                                                                          GENERAL INFORMATION:
APPLICANT: Yuan, Zhi-Min
APPLICANT: Gu, JiJie
TITLE OF INVENTION: Inhibition of p53 Degradation
FILE REFERENCE: 21508-044
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
Matches 18; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: PatentIn Ver. 2.1 SEQ ID NO 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 6, Application US/10397635
Publication No. US20040009604A1
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: POTENT ONCOLYTIC HERPES SIMPLEX VIRUS FOR CANCER TITLE OF INVENTION: THERAPY
FILE REFERENCE: P02410US2
CURRENT APPLICATION NUMBER: US/10/397,635
CURRENT FILING DATE: 2003-03-26
PRIOR APPLICATION NUMBER: 60/367,788
PRIOR FILING DATE: 2002-03-27
PRIOR FILING DATE: 2002-03-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: 60/410,024 PRIOR FILING DATE: 2002-09-11
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APPLICANT: FU, X
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQ ID NOS: 14
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INFORMATION FOR SEQ ID NO: 22:
SEQUENCE CHARACTERISTICS:
                                                       ENGTH: 1307
                                                                                                                                                                                                                                                                                                                                                                                                                             113 CCCGGAAGGCAGTCTGGC 96
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            35 CCCGGAAGGCAGTCTGGC 18
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TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FU, XUPING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%; Score 18; DB ilarity 100.0%; Pred. No. 7.3; Conservative 0; Mismatches
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RESULT 7
US-10-160-290-1/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: FastSEQ for Windows Version 3.0 SEQ ID NO 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
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Best Local
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PRIOR TILING DATE: 1997-12-04
PRIOR APPLICATION NUMBER: pctu896/01535
PRIOR FILING DATE: 1996-02-16
PRIOR APPLICATION NUMBER: 08/392,542
PRIOR FILING DATE: 1995-02-16
NUMBER OF SEQ ID NOS: 35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Halazonetis, Thanos
APPLICANT: Hartwig, Wolfgang
TITLE OF INVENTION: Peptides and peptidomimetics with
TITLE OF INVENTION: Structural similarity to human p53 that activate p53
TITLE OF INVENTION: function
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILE REFERENCE: 2973.1998
CURRENT APPLICATION NUMBER: US/09/829,922
CURRENT FILING DATE: 2001-04-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ENGTH: 1317
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
                                            COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

COMPUTER: IBM PC-DOS/MS-DOS

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Halazonetis, Thanos
Hartwig, Wolfgang
TITLE OF INVENTION: Peptides nad Peptidomimetics with
Signal Similarity to Human p53 That Activate
                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSEE: Banner,
                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     127
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18; Conservative
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        CLASSIFICATION: <Unknown>
                       APPLICATION NUMBER: US/10/160,290 FILING DATE: 04-Jun-2002
                                                                                                                                                                                                          COUNTRY: U.S.
                                                                                                                                                                                                                              STATE: District of Columbia
                                                                                                                                                                                                                                               CITY: Washington,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%; Silarity 100.0%; FC Conservative 0;
                                                                                                                                                                                                                                                 1001 G Street, N.W. ashington, D.C.
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Pred. No. 7
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                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Stockert, Elisabeth
APPLICANT: Chen, Yao-Tseng
TITLE OF INVENTION: Colon Cancer Antigen Panel
FILE REFERENCE: L0461/7105 (JRV)
CURRENT FILING DATE: 2001-05-04
NUMBER OF SEQ ID NOS: 30
SOFTWARE: PatentIn version 3.0
SEQ ID NO 9
LENGTH: 1760
TYPE: DNA
ORGANISM: Homo sapien
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                                                                                                        GENERAL INFORMATION:
                                                                                                                       Sequence 4, Application US/10191121
Publication No. US20040005574A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 9, Application US/09849602
Publication No. US20030165834A1
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Scanlan, Matthew J. APPLICANT: Old, Lloyd J. APPLICANT: Stockert, Elisabet
                                        APPLICANT: Guarente, Leonard
APPLICANT: Vaziri, Homayoun
APPLICANT: Imai, Shin-Ichiro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPHONE: 202 508-9100
TELEFAX: 202 508-9299
INFORMATION FOR SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                                                                                   Local
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CANT: Gu, Wei
OF INVENTION: Sir2 and p53 Deacetylation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ORGANISM: Homo sapiens SEQUENCE DESCRIPTION: SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY: linear MOLECULE TYPE: cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  REGISTRATION NUMBER: 34,698
REFERENCE/DOCKET NUMBER: 0486.48439
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE: <Unknown>
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                 206 CCCGGAAGGCAGTCTGGC 189
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                                                                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                                                     cccedaaeccaercreec 18
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FILING DATE: 10-Oct-2000
APPLICATION NUMBER: 08/392,542
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Posorske, Laurence H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ENGTH: 1317 base pairs
                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                               100.0%; Score 18; DB 10; 100.0%; Pred. No. 7;
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                                                                                                                                                                                                                                                                                                                            0;
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                                                                                                                                                                                                                                                                                                                                                                Length 1760;
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PRIOR APPLICATION NUMBER: US 60/303,456
PRIOR FILING DATE: 2001-07-06
PRIOR PELICATION NUMBER: US 10/NNN,NNN
PRIOR FILING DATE: 2002-07-05
PRIOR APPLICATION NUMBER: US 60/303,370
PRIOR FILING DATE: 2001-07-06
NUMBER OF SEQ ID NOS: 18
SOFTWARE: PATENTIN VERSION 3.1
SEQ ID NO 4
                                                                                                                                                                                                                                                          RESULT 11
US-10-429-802-25/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; NAME/KEY: misc_feature
; OTHER INFORMATION: Incyte ID No. US20040005559A1 430224.2
US-10-062-674-2065
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US-10-062-674-2065/c
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Publication No. US20040005559A1

GENERAL INFORMATION:

APPLICANT: Loring, Jeanne F.; Kaser, Matthew R.

TITLE OF INVENTION: MARKERS OF NEURONAL DIFFERENTIATION AND MORPHOGENESIS

FILE REFERENCE: PA-0026-1 CIP

CURRENT APPLICATION NUMBER: US/10/062,674

CURRENT FILING DATE: 2002-01-30
                                                                                                                                                                                            Sequence 25, Application US/10429802
publication No. US20030228285A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: US 09/625,102
PRIOR FILING DATE: 2000-07-24
NUMBER OF SEQ ID NOS: 2217
SOFTWARE: PERL Program
SEQ ID NO 2065
LENGTH: 2209
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Best Local Similarity
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APPLICANT: HUNG, MIEN-CHIE
APPLICANT: WONG, KA YIN
APPLICANT: WONG, KA YIN
APPLICANT: ZOU, YIYU
TITLE OF INVENTION: BIPARTITE T-CELL FACTOR (TCF)-RESPONSIVE PROMOTER
FILE REFERENCE: UTSC:752US
CURRENT PELICATION UNUMERR: US/10/429,802
CURRENT FILING DATE: 2003-05-05
PRIOR APPLICATION NUMBER: 60/377,672
PRIOR FILING DATE: 2002-05-03
NUMBER OF SEQ ID NOS: 54
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CURRENT APPLICATION NUMBER: US/10/191,121
CURRENT FILING DATE: 2002-07-08
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ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Homo sapiens FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: DNA
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ilarity 100.0%;
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Pred. No.
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6.8;
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APPLICANT: HUNG, MIEN-CHIE
APPLICANT: LAW, KENG-LI
APPLICANT: LOYANG, FU
APPLICANT: LIU, JAW-CHING
APPLICANT: LIU, JAW-CHING
APPLICANT: LIU, JAW-CHING
TITLE OF INVENTION: TARGETING PROTEINS TO DELIVER THERAPEUTIC OR DIAGNOSTIC
TITLE OF INVENTION: REAGENTS
TILLE OF INVENTION NUMBER: US/10/430,503
CURRENT APPLICATION NUMBER: US/10/430,503
CURRENT FILING DATE: 2003-05-06
PRIOR APPLICATION NUMBER: 60/383,063
PRIOR APPLICATION NUMBER: 60/383,063
PRIOR APPLICATION NUMBER: 60/383,063
PRIOR APPLICATION PRIOR : 2002-05-06
NUMBER OF SEQ ID NOS: 42
SOFTWARE: Patentin Ver: 2.1
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US-10-391-068-1/c
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; ORGANISM: Homo sapiens
US-10-430-503-16
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                                                                                                                                                                                                                                                                                                                                                                     Sequence 1, Application US/10391068
Publication No. US20040028654A1
GENERAL INFORMATION:
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SEQ ID NO 1
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SEQ ID NO 25
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Best Local
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APPLICANT: ROTH, JACK
APPLICANT: ROTH, JACK
TITLE OF INVENTION: PROTAMINE-ADENOVIRAL VECTOR COMPLEXES
TITLE OF INVENTION: AND METHODS OF USE
FILE REFERENCE: INRP:097US
CURRENT APPLICATION NUMBER: US/10/391,068
CURRENT FILING DATE: 2003-03-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity
                                                                                                                                                                  PRIOR APPLICATION NUMBER: 60/366,846 PRIOR FILING DATE: 2002-03-22 NUMBER OF SEQ ID NOS: 32
NAME/KEY: CDS
LOCATION: (252)..(1433)
                                                            TYPE: DNA
ORGANISM: Homo sapiens
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                                                                                                     ENGTH:
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18; Conserv
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RESULT 15
US-10-077-176-59/c
; Sequence 59, Application US/10077176
; Publication No. US20030175862A1
; GENERAL INFORMATION:
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US-10-077-176-58/c
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CURRENT FILING NUMBER: US/10/077,176
CURRENT FILING DATE: 2002-02-19
NUMBER OF SEQ ID NOS: 71
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 58
LENGTH: 2629
                                                    Sequence 60, Application US/10077176 Fublication No. US20030175862A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/10/077,176
CURRENT FILING DATE: 2002-02-19
NUMBER OF SEQ ID NOS: 71
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 59
LENGTH: 2629
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Best Local Similarity
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Best Local :
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TITLE OF INVENTION: ENGINEERED OPEN READING FRAME FOR
FILE REFERENCE: 004255.00008
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TITLE OF INVENTION: ENGINEERED OPEN READING FRAME FOR P53
FILE REFERENCE: 004255.00008
APPLICANT: Brachmann, Rainer
TITLE OF INVENTION: ENGINEERED OPEN READING FRAME FOR P53
FILE REFERENCE: 004255.00008
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ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                     TYPE: DNA ORGANISM: Homo sapiens
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100.0%; Pred. No. 6.7;
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Pred. No. 6.
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6.7;
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; LENGTH: 2629
; TYPE: DNA
; ORGANISM: Homo s
US-10-07,7-176-61
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US-10-077-176-61/c
; Sequence 61, Application US/10077176
; Publication No. US20030175862A1
; GENERAL INFORMATION:
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                                                                   ; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-439-388-16
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US-10-439-388-16/c
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CURRENT FILING DATE: 2002-02-19
NUMBER OF SEQ ID NOS: 71
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 60
                                                                                                                                  CURRENT APPLICATION NUMBER: US/10/439,388
CURRENT FILING DATE: 2003-05-16
PRIOR APPLICATION NUMBER: US 60/381,055
PRIOR FILING DATE: 2002-05-16
NUMBER OF SEQ ID NOS: 70
SOFTWARE: PatentIn version 3.2
SEQ ID NO 16
SEQ ID NO 16
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SOFTMARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 61
LENGTH: 2629
                                                                                                                                                                                                                                                                                                                           Sequence 16, Application US/10439388 Publication No. US20030228617A1 GENERAL INFORMATION:
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                  Query Match
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TITLE OF INVENTION: ENGINEERED OPEN READING
FILE REFERENCE: 004255, 00008
CURRENT APPLICATION NUMBER: US/10/077,176
CURRENT FILING DATE: 2002-02-19
                                                                                                                                                                                                                                                      APPLICANT: Aune, Thomas M
APPLICANT: Olsen, Nancy J
TITLE OF INVENTION: Method for Predicting
FILE REFERENCE: 1242/68
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TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                      ENGTH:
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18; Conserv
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100.0%; Score 18; I larity 100.0%; Pred. No. 6. Conservative 0; Mismatches
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CCCGGAAGGCAGTCTGGC 18

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CCCGGAAGGCAGTCTGGC 226

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RESULT 21
US-09-918-995-36880/c
Sequence 36880, Application US/09918995
Publication No. US20030073623A1
RENERAL INFORMATION:
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US-10-159-563-188/c
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US-10-165-216-3/c
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Publication No. US20030228675A1
GENERAL INFORMATION:
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LENGTH: 2629
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APPLICANT: Abraham, Robert T.
TITLE OF INVENTION: ATM Related Kinase ATX, Nucleic Acids
TITLE OF INVENTION: Encoding Same and Methods of Use
TILE REFERENCE: P-LJ 5222
CURRENT APPLICATION NUMBER: US/10/165,216
CURRENT FILING DATE: 2002-06-06
NUMBER OF SEQ ID NOS: 11
                                                                                                                                                                                                                                                                        Query Match
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SEQ ID NO 188
LENGTH: 2629
                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Peterson, Carsten
APPLICANT: Meltzer, Paul
APPLICANT: Meltzer, Paul
TITLE OF INVENTION: SELECTIONS OF GENES AND METHODS OF USING THE SAME FOR
TITLE OF INVENTION: DIAGNOSIS AND FOR TARGETING THE THERAPY OF SELECT CANCERS
FILE REFERENCE: 11613.56USI1
CURRENT PEPLICATION NUMBER: US/10/159,563
CURRENT FILING DATE: 2002-12-09
PRIOR APPLICATION NUMBER: US 10/133,937
PRIOR PERIOR DATE: 2002-04-25
PRIOR FILING DATE: 2002-04-25
NUMBER OF SEQ ID NOS: 444
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Khan, Javed
APPLICANT: Ringner, Markus
APPLICANT: Peterson, Carste
APPLICANT: Meltzer, Paul
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ORGANISM: Homo sapiens
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LOCATION: (252)...(1433)
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Pred. No.
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APPLICANT: Hyseq, Inc.
TITLE OF INVENTION: NOVEL NUCLEIC ACID SEQUENCES OBT
TITLE OF INVENTION: FROM VARIOUS CDNA LIBRARIES
FILE REFERENCE: 20411-756
CURRENT APPLICATION NUMBER: US/09/918,995
CURRENT FILING DATE: 2001-07-30
PRIOR APPLICATION NUMBER: US/09/235,076
PRIOR APPLICATION NUMBER: US/09/235,076
PRIOR FILING DATE: 1999-01-20
NUMBER OF FILING DATE: 1999-01-20
NUMBER OF SEQ ID NOS: 38054
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 36880
LENGTH: 436
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                                                                                                                                                 US-10-066-543-113
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CURRENT FILING DATE: 2002-01-31
NUMBER OF SEQ ID NOS: 3417
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 113
LENGTH: 905
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Best Local :
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                                                                                                                                                     TYPE: DNA

ORGANISM: Homo sapiens

PEATURE:

NAME/KEY: misc feature

NAME/KEY: misc feature

LOCATION: 610, 616, 621, 623, 624, 625, 629, 630, 646, 652, 670, 672, 600

LOCATION: 674, 676, 688, 700, 709, 710, 744, 753, 756, 758, 759, 763, 100

LOCATION: 786, 796, 823, 851, 876, 885

OTHER INFORMATION: n = A,T,C or G
                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY TITLE OF INVENTION: AND DIAGNOSIS OF COLON CANCER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               10-066-543-113/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                FILE REFERENCE: 210121.563
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Jiang, Yugiu
APPLICANT: Pyle, Ruth A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
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blication No. US20030087818A1
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 516 CCCAGAAGGCAGTCTGGC
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17; Conserv
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Carter, Darrick
Fanger, Gary R.
Smith, Carole L.
Durham, Margarita
Stolk, John A.
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94.4%; Pred. No. 51;
ive 0; Mismatches
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Pred. No. 56;
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RESULT 23 US-09-764-847-1845/c ; Sequence 1845, Application US/09764847

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RESULT 25
US-10-092-154-1845/c
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US-09-764-847-1845
                                  US-10-092-154-1845
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GENERAL INFORMATION:
APPLICANT: Rosen et al.
TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
FILE REFERENCE: PCO09
CURRENT APPLICATION NUMBER: US/09/764,847
CURRENT FILING DATE: 2001-01-17
                                                                                             CURRENT APPLICATION NUMBER: US/10/092,154
CURRENT FILING DATE: 2002-03-07
NUMBER OF SEQ ID NOS: 2003
Prior Application removed - See File Wrapper or Palm SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 1845
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
APPLICANT: Rosen et al.
APPLICANT: Rosen et al.
TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
FILE REFERENCE: PC009
CURRENT APPLICATION NUMBER: US/09/764,847
CURRENT FILING DATE: 2001-01-17
Prior application data removed - consult PALM or file wrapper
NUMBER OF SEQ ID NOS: 2003
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 1846
LENGTH: 942
                                                                                                                                                                                                  Sequence 1845, Application US/10092154
Publication No. US20030054375A1
GENERAL INFORMATION:
APPLICANT: Rosen et al.
TITLE OF INVENTION: Nucleic Acids, Proteins,
FILE REFERENCE: PC009C1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQ ID NO 1845
LENGTH: 942
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Query Match
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                                             LENGTH: 942
TYPE: DNA
ORGANISM: Homo:
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Pred. No. 50;
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Length 942;
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US-09-765-231A-2/c
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                                                                                                                     LENGTH: 1986
TYPE: DNA
ORGANISM: Homo sapiens
US-09-765-231A-2
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SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 1846
LENGTH: 942
TYPE: DNA
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GENERAL INFORMATION:
APPLICANT: Rosen et
                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/09/765,231A
CURRENT FILING DATE: 2001-01-18
NUMBER OF SEQ ID NOS: 82
SEQ ID NO 2
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Matches 17; Conserv
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Best Local 9
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Patent No. US20020119452A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/10/092,154
CURRENT FILING DATE: 2002-03-07
NUMBER OF SEQ ID NOS: 2003
                                                                                                                                                                                                                                                                                      APPLICANT: Vasanthakamu
APPLICANT: Dotson, Stant
APPLICANT: Ma, Xiao-Jun
                                                                                                                                                                                                                                                                                                                                   APPLICANT: Searle/Monsanto APPLICANT: Phippard, Deborah
                                                                                                                                                                                                                                          TITLE OF INVENTION: Osteoarthritis tissue-derived nucleic acids, polypeptides, TITLE OF INVENTION: vectors, and cells FILE REFERENCE: SO-3221 PR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TIE OF INVENTION: Nucleic Acids, Proteins, and Antibodies LE REFERENCE: PC009C1
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893 CCCAGAAGGCAGTCTGGC 876
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17; Conserv
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Dotson, Stanton
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Conservative
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94.4%;
                                                                         91.1%; Score 16.4;
94.4%; Pred. No. 46;
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Pred. No. 5
                                                            Mismatches
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RESULT 28 US-09-884-441-332

Sequence 332, Application US/09884441
Patent No. US20020119158A1
GENERAL INFORMATION:

APPLICANT: Algate, Paul A.

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CURRENT APPLICATION NUMBER: US/09/884,441
CURRENT FILING DATE: 2001-06-18
NUMBER OF SEQ ID NOS: 489
SOFTWARE: FRATSEQ for Windows Version 3.0
SEQ ID NO 332
LENGTH: 184
GENERAL INFORMATION:

APPLICANT: Retter, Marc W.

APPLICANT: Fanger, Gary R.

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND TITLE OF INVENTION: DIAGNOSIS OF OVARIAN CANCER FILE REFERENCE: 210121.462C6

CURRENT APPLICATION NUMBER: US/09/827,271

CURRENT FILING DATE: 2001-04-04
                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQ ID NO 332
LENGTH: 184
                                                                                                                                            Sequence 332, A Publication No.
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TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.462C8
CURRENT APPLICATION NUMBER: US/09/907,969
CURRENT FILING DATE: 2001-07-17
NUMBER OF SEQ ID NOS: 596
SOFTWARE: FastSEQ for Windows Version 4.0
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TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR
TITLE OF INVENTION: DIAGNOSIS OF OVARIAN CANCER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Mitcham
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Algate, Paul A.
Fling, Steven P.
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Vedvick, Thomas S.
Carter, Darrick
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Fanger, Gary Richard
                                                                                                                                            Application US/09827271 o. US20030165504A1
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94.1%;
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Pred. No. 2
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Pred. No. 2e+02;
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APPLICANT: Chondrogene Inc.
APPLICANT: Liew, C.C.
APPLICANT: Liew, C.C.
APPLICANT: Liew, C.C.
FILE REFERENCE: 4231/2002
CURRENT APPLICATION NUMBER: US/10/085,783A
CURRENT APPLICATION NUMBER: US/07/085,783A
CURRENT FILING DATE: 2002-02-28
PRIOR APPLICATION NUMBER: US/07/5,017
PRIOR FILING DATE: 2001-07-13
PRIOR FILING DATE: 2001-03-12
PRIOR APPLICATION NUMBER: US/07/5,017
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US-10-198-053-332
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US-10-198-053-332
; ORGANISM: Human
US-10-085-783A-47890
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 332
LENGTH: 184
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Best Local :
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SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 332
LENGTH: 184
                                                                                   SOFTWARE: PatentIn version 3.2 SEQ ID NO 47890
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Publication No.
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CURRENT APPLICATION NUMBER: US/10/198,053
CURRENT FILING DATE: 2002-07-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Retter, Marc W.
APPLICANT: Ranger, Gary R.
APPLICANT: Hill, Paul
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Bangur, Chaitanya S. APPLICANT: Retter, Marc W.
                                                                                                                            NUMBER OF SEQ ID NOS: 58994
                                                                ENGTH: 358
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Local Similarity 94.1%;
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b. US20030124140A1
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94.1%;
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Pred. No. 2
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Pred. No. 2
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US-10-242-535A-47890/c
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Best Local S
Matches 16
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                                                                                                                               SEQ ID NO 21425
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PRIOR APPLICATION NUMBER: US 10/085,783
PRIOR FILING DATE: 2002-02-28
PRIOR APPLICATION NUMBER: US 60/305,340
PRIOR FILING DATE: 2001-07-13
                                                                                                                                                           PRIOR APPLICATION NUMBER: US 60/275,017
PRIOR FILING DATE: 2001-03-12
PRIOR APPLICATION NUMBER: US 60/271,955
PRIOR FILING DATE: 2001-02-28
NUMBER OF SEQ ID NOS: 58994
                                                                                                                                                                                                                                                        FILE REFERENCE: 4231/2002
CURRENT APPLICATION NUMBER: US/10/085,783A
CURRENT FILING DATE: 2002-02-28
PRIOR APPLICATION NUMBER: US 60/305,340
PRIOR FILING DATE: 2001-07-13
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EQ ID NO 47890
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PRIOR FILING DATE: 2001-03-12
PRIOR APPLICATION NUMBER: US 60/271,955
PRIOR FILING DATE: 2001-02-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/10/242,535A CURRENT FILING DATE: 2002-09-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: ChondroGene Inc. APPLICANT: Liew, C.C.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQ ID NOS: 58994
                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: ChondroGene Inc. APPLICANT: Liew, C.C.
                                                                                                                                               SOFTWARE: PatentIn version 3.2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 358
TYPE: DNA
ORGANISM: Human
                   NAME/KEY: misc_feature
LOCATION: (1)..(1)
                                                       ORGANISM: Human FEATURE:
OTHER INFORMATION: n is a,
                                                                                                            ENGTH: 371
                                                                                                                                                                                                                                                                                                                                                    TLE OF INVENTION: Compositions and Methods Relatiing to Osteoarthritis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           358 CCAGAAGGCAGTCTGGC 342
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Pred. No. 1
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Pred. No. 1
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TITLE OF INVENTION: Compositions and Methods Relating to FILE REFERENCE: 4231/2005
CURRENT APPLICATION NUMBER: US/10/242,535A
CURRENT FILING DATE: 2002-09-12
PRIOR APPLICATION NUMBER: US 10/085,783
PRIOR APPLICATION NUMBER: US 60/305,340
PRIOR PILING DATE: 2001-07-13
PRIOR FILING DATE: 2001-07-13
PRIOR APPLICATION NUMBER: US 60/275,017
PRIOR APPLICATION NUMBER: US 60/275,955
PRIOR APPLICATION NUMBER: US 60/271,955
PRIOR FILING DATE: 2001-03-12
PRIOR FILING DATE: 2001-03-12
PRIOR FILING DATE: 2001-03-12
PRIOR FILING DATE: 2001-03-12
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; OTHER INFORMATION: n is
US-10-085-783A-21425
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                                                                                                                                                                                                                                                                                                                                                                 SEQ ID NO 21425
LENGTH: 371
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Best Local Similarity
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                                                                                          Query Match
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SOFTWARE: PatentIn ver
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ublication No. US20040013663A1
ENERAL INFORMATION:
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LOCATION: (205)...(206)
OTHER INFORMATION: n is a,
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LOCATION: (203)..(203)
OTHER INFORMATION: n is
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NAME/KEY: misc_feature
LOCATION: (1)..(1)
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165 CCAGAAGGCAGTCTGGC 149
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                            2 CCGGAAGGCAGTCTGGC 18
                                                                            Similarity
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Pred. No. 1.
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                                                                                           Length 371;
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US-10-085-783A-9459/c
; Sequence 9459, Application US/10085783A
; Publication No. US20040037841A1

GENERAL INFORMATION:
APPLICANT: ChondroGene Inc.
APPLICANT: Liew, C.C.

RESULT 36

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FILE REFERENCE: 4231/2002

CURRENT APPLICATION NUMBER: US/10/085,783A

CURRENT FILING DATE: 2002-02-28

PRIOR APPLICATION NUMBER: US 60/305,340

PRIOR FILING DATE: 2001-07-13

PRIOR APPLICATION NUMBER: US 60/275,017

PRIOR FILING DATE: 2001-03-12

PRIOR APPLICATION NUMBER: US 60/271,955

PRIOR APPLICATION NUMBER: US 60/271,955

PRIOR FILING DATE: 2001-02-28

NUMBER OF 5EQ ID NOS: 58994

SOFTWARE: Patentin version 3.2

SEQ ID NO 9459

LENGTH: 395
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; LOCATION: (63)...(63)
; OTHER INFORMATION: n is a,
US-10-085-783A-9459
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 9459, Application US/10242535A Publication No. US20040013663A1
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LENGTH: 395
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                                                                                                                                                                                                                                                                                                   APPLICANT: Liew, C.C. Compositions and Methods Relating to Osteoarthritis FILE REFERENCE: 4231/2005

CURRENT APPLICATION NUMBER: US/10/242,535A

CURRENT FILING DATE: 2002-09-12

PRIOR APPLICATION NUMBER: US 10/085,783

PRIOR FILING DATE: 2002-02-28

PRIOR PPLICATION NUMBER: US 60/305,340

PRIOR APPLICATION NUMBER: US 60/305,340

PRIOR FILING DATE: 2001-07-13

PRIOR PILING DATE: 2001-07-13

PRIOR PILING DATE: 2001-03-12

PRIOR APPLICATION NUMBER: US 60/275,017

PRIOR PILING DATE: 2001-03-12

PRIOR PILING DATE: 2001-03-12

PRIOR PILING DATE: 2001-03-12

PRIOR PILING DATE: 2001-03-28
                                                                                                                                                                                                                                                            PRIOR FILING DATE: 2001-02-28
NUMBER OF SEQ ID NOS: 58994
SOFTWARE: PatentIn version 3.2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: ChondroGene Inc. APPLICANT: Liew, C.C.
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LOCATION: (21)..(22)
OTHER INFORMATION: n is a,
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OTHER INFORMATION: n is
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ORGANISM: Human
                       NAME/KEY: misc_feature
LOCATION: (36)..(36)
                                                                                   NAME/KEY: misc feature LOCATION: (21)...(22) OTHER INFORMATION: n is
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  INFORMATION: n is
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Pred. No. 1.8e+02;
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; LOCATION: (212)...(212)
; OTHER INFORMATION: n is
US-10-085-783A-23153
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; LCCATION: (63)..(63)
; OTHER INFORMATION: n is a,
US-10-242-535A-9459
                                                                                                                                                                                               RESULT 39
US-10-242-535A-23153/c
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                                                                                                                                         Sequence 23153, Application US/10242535A Publication No. US20040013663A1 GENERAL INFORMATION:
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Best Local (
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APPLICANT: Chondrogene Inc.
APPLICANT: Liew, C.C.
ITTLE OF INVENTION: Compositions and Methods Relating
FILE REFERENCE: 4231/2005
CURRENT APPLICATION NUMBER: US/10/242,535A
CURRENT FILING DATE: 2002-09-12
PRIOR APPLICATION NUMBER: US 10/085,783
PRIOR FILING DATE: 2002-02-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Liew, C.C.
TITLE OF INVENTION: Compositions and Methods Relating to Osteoarthritis
FILE REFERENCE: 4231/2002
CURRENT APPLICATION NUMBER: US/10/085,783A
CURRENT FILING DATE: 2002-02-28
PRIOR APPLICATION NUMBER: US 60/305,340
PRIOR FILING DATE: 2001-07-13
PRIOR APPLICATION NUMBER: US 60/375,017
PRIOR APPLICATION NUMBER: US 60/275,017
PRIOR APPLICATION NUMBER: US 60/275,017
PRIOR APPLICATION NUMBER: US 60/275,017
PRIOR FILING DATE: 2001-03-12
PRIOR FILING DATE: 2001-03-12
PRIOR FILING DATE: 2001-02-28
NUMBER OF SEQ ID NOS: 58994
COUNTRIEN DESCRIPT OF SEG 12
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APPLICANT: Liew, C.C.
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OTHER INFORMATION: n is
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LOCATION: (56)..(56)
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ORGANISM: Human
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Pred. No. 1.8e
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1.8e+02;
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                                                                                          to Osteoarthritis
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CURRENT APPLICATION NUMBER: US/10/085,783A

CURRENT FILING DATE: 2002-02-28

PRIOR APPLICATION NUMBER: US 60/305,340

PRIOR FILING DATE: 2001-07-13

PRIOR APPLICATION NUMBER: US 60/275,017

PRIOR APPLICATION NUMBER: US 60/271,955

PRIOR APPLICATION NUMBER: US 60/271,955

PRIOR FILING DATE: 2001-02-28

PRIOR FILING DATE: 2001-02-28
                                                                               Query Match
Best Local :
                                                               Matches
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TYPE: DNA
ORGANISM: Human
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APPLICANT: Liew, C.C.
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                                                                                                                                           NAME/KEY: misc_feature
LOCATION: (42)...(42)
OTHER INFORMATION: n is a,
                                                                                                                                                                                                              TYPE: DNA
ORGANISM: Human
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LOCATION: (212)..(212)
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OTHER INFORMATION: n is
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JOCATION: (67)..(68)
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OCATION: (56)..(56)
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351 CCAGAAGGCAGTCTGGC 335
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LING DATE: 2001-02-28
                                                               l Similarity
16; Conserv
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Similarity 94.1%;
                             CCGGAAGGCAGTCTGGC 18
                                                               Conservative
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Pred. No. 1.8e+02;
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Pred. No. 1.8e+02;
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US-10-085-783A-23133/c
; Sequence 23133, Application US/10085783A
; Publication No. US20040037841A1
; GENERAL INFORMATION:
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; OTHER INFORMATION: n is
US-10-242-535A-23042
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SEQ ID NO 23042
LENGTH: 399
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APPLICANT: Liew, C.C.
TITLE OF INVENTION: Compositions and Methods Relating to Osteoarthritis
FILE REFERENCE: 4231/2005
FILE REFERENCE: 4231/2005
CURRENT FALING DATE: 2002-09-12
PRIOR APPLICATION NUMBER: US/10/242,535A
CURRENT FILING DATE: 2002-09-12
PRIOR APPLICATION NUMBER: US 10/085,783
PRIOR FILING DATE: 2002-02-28
PRIOR APPLICATION NUMBER: US 60/305,340
PRIOR FILING DATE: 2001-07-13
PRIOR FILING DATE: 2001-07-13
PRIOR APPLICATION NUMBER: US 60/275,017
PRIOR APPLICATION NUMBER: US 60/275,017
PRIOR APPLICATION NUMBER: US 60/271,955
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PRIOR FILING DATE: 2001-07-13
PRIOR PPLICATION NUMBER: US 60/275,017
PRIOR FILING DATE: 2001-03-12
PRIOR APPLICATION NUMBER: US 60/271,955
PRIOR PILING DATE: 2001-02-28
NUMBER FILING DATE: 2001-02-28
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ORGANISM: Human
FEATURE:
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CURRENT FILING DATE: 2002-02-28
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APPLICANT: Liew, C.C.

IITLE OF INVENTION: Compositions and Methods Relatiing to Osteoarthritis

FILE REFERENCE: 4231/2002
                                                             FEATURE:
NAME/KEY: misc_feature
LOCATION: (40)..(40)
NAME/KEY: misc_feature
                                                                                                                                                                                       ORGANISM: Human
FEATURE:
                                                                                                                                                                                                                            TYPE: DNA
                                                                                                                         OTHER INFORMATION: n is
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                                            OTHER INFORMATION: n is
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NAME/KEY: misc feature LOCATION: (341)...(341) OTHER INFORMATION: n is a,

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LOCATION: (48)..(48)
OTHER INFORMATION: n

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SEQ ID NO 23133
LENGTH: 400.
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Matches
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                               Matches
                                                           Query Match
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PRIOR FILING DATE: 2002-02-28
PRIOR APPLICATION NUMBER: US 60/305,340
PRIOR FILING DATE: 2001-07-13
PRIOR APPLICATION NUMBER: US 60/275,017
PRIOR FILING DATE: 2001-03-12
PRIOR APPLICATION NUMBER: US 60/271,955
PRIOR APPLICATION NUMBER: US 60/271,955
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APPLICANT: Liew, C.C.
TITLE OF INVENTION: Compositions and Methods Relatiing to Osteoarthritis
FILE REFERENCE: 4231/2005
CURRENT APPLICATION NUMBER: US/10/242,535A
CURRENT FILING DATE: 2002-09-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR FILING DATE: 2001-02-28
NUMBER OF SEQ ID NOS: 58994
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LOCATION: (394)..(394)
OTHER INFORMATION: n is a,
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LOCATION: (341)..(341)
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LOCATION: (40)..(40)
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LOCATION: (7)..(7)
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16; Conservative
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CURRENT FILING DATE: 2001-08-03

NUMBER OF SEQ ID NOS: 1124

SOFTWARE: FASTSEQ for Windows Version 4.0

SEQ ID NO 755

LENGTH: 405

TYPE: DNA

ORGANISM: Homo sapiens
FEATURE:
NAME/KEN: 214, 305, 366, 368, 378

OTHER INFORMATION: n = A,T,C or G

US-09-922-217-755
OTHER INFORMATION: n = A,T,C or US-09-833-263-755
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US-09-922-217-755/c
                                                                                                                                         FILE REFERENCE: 210121.471C12
CURRENT APPLICATION NUMBER: US/09/833,263
CURRENT FILING DATE: 2001-04-10
NUMBER OF SEQ ID NOS: 1093
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 755
                                                                                                                                                                                                                                                                                                                                                       Patent No. US2002011 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS TITLE OF INVENTION: OF COLON CANCER AND METHODS FOR THEIR USE FILE REFERENCE: 210121.471C13
CURRENT APPLICATION NUMBER: US/09/922,217
CURRENT FILING DATE: 2001-08-03
NUMBER OF SEC. TO THE TIME DATE: 2001-08-03
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Best Local :
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APPLICANT: Clapper, Jonat
APPLICANT: Stolk, John A.
APPLICANT: Meagher, Madel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
                                                                                                                                                                                                                                              APPLICANT: Meagher, Madeleine J.
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF COLON CANCER AND METHODS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT
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                                        FEATURE:
NAME/KEY: misc_feature
LOCATION: (1)...(405)
                                                                                        LENGTH: 405
TYPE: DNA
ORGANISM: Homo sapien
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16; Conserv
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Benson, Darin R.
Meagher, Madeleine J
Stolk, John A.
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Smith, Carole Lynn
King, Gordon E.
Wang, Aijun
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1.8e+02;
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; FEATURE:

NAME/KEY: misc feature

; LOCATION: 214, 305, 366, 368,

; OTHER INFORMATION: n = A,T,C or

US-10-025-380-755
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US-10-025-380-755/c
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                                                                                                                                                                                                             Sequence 58222, Application US/10085783A Publication No. US20040037841A1 GENERAL INFORMATION:
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Best Local
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                                                                                                   APPLICANT: Chondrogene Inc.
APPLICANT: Liew, C.C.
TITLE OF INVENTION: Compositions and Methods Relating
FILE REFERENCE: 4231/2002
CURRENT APPLICATION NUMBER: US/10/085,783A
CURRENT FILING DATE: 2002-02-28
CURRENT FILING DATE: 2002-02-28
     PRIOR APPLICATION NUMBER: US 60/305,340 PRIOR FILING DATE: 2001-07-13 PRIOR APPLICATION NUMBER: US 60/275,017 PRIOR FILING DATE: 2001-03-12 PRIOR APPLICATION NUMBER: US 60/271,955 PRIOR FILING DATE: 2001-02-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/10/025,380
CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 1129
SOFTWARE: FastSEQ for Windows Version 4.0
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TYPE: DNA
ORGANISM: Homo sapiens
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LE OF INVENTION: OF COLON CANCER AND METHODS FOR THEIR USE
E REFERENCE: 210121.471C14
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Benson, Darin R.
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94.1%;
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Pred. No. 1
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.8e+02;
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Sequence 17255, Application US/10085783A
Publication No. US20040037841A1
GENERAL INFORMATION:
APPLICANT: ChondroGene Inc.
APPLICANT: Liew, C.C.
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Matches
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SEQ ID NO 58222
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Publication No.
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APPLICANT: Liew, C.C.
TITLE OF INVENTION: Compositions and Methods Relatiing to Osteoarthritis
FILE REFERENCE: 4231/2002
CURRENT APPLICATION NUMBER: US/10/085,783A
CURRENT FILING DATE: 2002-02-28
PRIOR APPLICATION NUMBER: US 60/305,340
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: DNA
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US-10-242-535A-58222
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SOFTWARE: PatentIn version 3.2
SEQ ID NO 58222
LENGTH: 422
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Liew, C.C.
TITLE OF INVENTION: Compositions and Methods Relating to Osteoarthritis
FILE REFERENCE: 4211/2005
CURRENT APPLICATION NUMBER: US/10/242,535A
CURRENT FILING DATE: 2002-09-12
PRIOR APPLICATION NUMBER: US 10/085,783
PRIOR FILING DATE: 2002-02-28
PRIOR FILING DATE: 2002-07-13
PRIOR PPLICATION NUMBER: US 60/305,340
PRIOR PPLICATION NUMBER: US 60/305,340
PRIOR PPLICATION NUMBER: US 60/275,017
PRIOR PILING DATE: 2001-03-12
PRIOR FILING DATE: 2001-03-12
PRIOR FILING DATE: 2001-03-12
                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: US 60/271,955
PRIOR FILING DATE: 2001-02-28
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FEATURE:
FATURE:
NAME/KEY: misc_feature
LOCATION: (56)...(56)
OTHER INFORMATION: n is a, c,
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                                                                                                                                                                                  NAME/KEY: misc_feature LOCATION: (56)..(56)
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                                                                                         Similarity
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CCAGAAGGCAGTCTGGC 289
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                                                                       Conservative
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94.1%;
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Pred. No. 1.8e+02;
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Pred. No. 1.
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GENERAL INFORMATION:
APPLICANT: Chondrogene Inc.
APPLICANT: Liew, C.C.
APPLICANT: Liew, C.C.
TITLE OF INVENTION: Compositions and Methods Relating to Osteoarthritis
FILE REFERENCE: 4231/2005
CURRENT PPLICATION NUMBER: US/10/242,535A
CURRENT FILING DATE: 2002-09-12
PRIOR APPLICATION NUMBER: US 10/085,783
PRIOR FILING DATE: 2001-07-13
PRIOR PILING DATE: 2001-07-13
PRIOR APPLICATION NUMBER: US 60/305,340
PRIOR FILING DATE: 2001-07-13
PRIOR RILING DATE: 2001-07-13
PRIOR RILING DATE: US-00/275,017
PRIOR APPLICATION NUMBER: US 60/275,917
PRIOR PILING DATE: 2001-03-12
PRIOR PILING DATE: 2001-03-12
PRIOR PILING DATE: 2001-03-28
NUMBER OF SEQ ID NOS: 58994
SOFTWARE: Patentin version 3.2
SEQ ID NO 17255
LENGTH: 443
TYPE: DNA
JORGANISM: Human
US-10-242-535A-17255
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PRIOR APPLICATION NUMBER: US 60/275,017
PRIOR FILING DATE: 2001-03-12
PRIOR APPLICATION NUMBER: US 60/271,955
PRIOR FILING DATE: 2001-02-28
PRIOR FILING DATE: 2001-02-28
VNUMBER OF SEQ ID NOS: 58994
SOPTWARE: Patentin version 3.2
SEQ ID NO 17255
LENGTH: 443
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US-10-242-535A-17255/c
US-10-242-535A-17255/c
; Sequence 17255, Application US/10242535A
; Publication NO. US20040013663A1
; GENERAL INFORMATION:
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Search completed: March 5, 2004, 02:19:51
Job time: 282 secs
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; ORGANISM: Human
US-10-085-783A-17255
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Best Local &
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Best Local Similarity
Matches 16; Conser
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Local Similarity 94.1%;
les 16; Conservative 0
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33 CCAGAAGGCAGTCTGGC 17
                                                                                      33 CCAGAAGGCAGTCTGGC 17
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                                                                                                                                                                        85.6%; Score 15.4; DB 15; Length 443; ilarity 94.1%; Pred. No. 1.8e+02; Conservative 0; Mismatches 1; Indels 0
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Pred. No. 1.8e+02;
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Sequence:

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2: /cgn2 6/ptcdata/2/ina/5B_COMB.seq:*
3: /cgn2 6/ptcdata/2/ina/6B_COMB.seq:*
4: /cgn2 6/ptcdata/2/ina/6B_COMB.seq:*
5: /cgn2 6/ptcdata/2/ina/6B_COMB.seq:*
6: /cgn2 6/ptcdata/2/ina/backfIles1.seq:*
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US-08-047-041A-15/c
Sequence 15, Application Uses No. 5527676
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APPLICANT: Nigro, Janice M.
ITILE OF INVENTION: Detection of Loss
TUTLE OF INVENTION: Gene
NUMBER OF SEQUENCES: 28
CORRESPONDENCE ADDRESS:
ADDRESSE: Banner & Allegretti, Ltd.
STREET: 1001 G Street, N.W.
CITY: Washington
STATE: D.C.
COUNTRY.
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                              STREET: 1001 G Street, N.W.

STREET: 1001 G Street, N.W.

CITY: Washington

STATE: D.C.

COUNTRY: USA
ZIP: 20001.4597

COMPUTER READABLE FORM.

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ver
COMPUTER: USATA:

OPERATION STREET: US/08/047,041A

FILING DATE: 22-MAR-1993

CLASSIFICATION NUMBER: US/09/28,661
APPLICATION NUMBER: US 07/928,661
APPLICATION NUMBER: US 07/928,661
APPLICATION NUMBER: US 07/446,584
APPLICATION NUMBER: US 07/446,584
APPLICATION NUMBER: US 07/330,566
APPLICATION NUMBER: US 07/330,566
FILING DATE: 29-MAR-1989
ATTORNEY/AGENT INFORMATION
AMME: KAGAN, SATAN A.

DESCRIPTANTON DATA:

APPLICATION NUMBER: US 07/330,566

FILING DATE: 29-MAR-1989
ATTORNEY/AGENT INFORMATION

NAME: KAGAN, SATAN A.
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181
182
NAME: Kagan, Sarah A.
REGISTRATION NUMBER: 32,141
REFERENCE/DOCKET NUMBER: 01
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US-09-371-696-1
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2490, App 2246, Appl 3246, Appl 3700, Appli 10014, Appli 10014, Appli 10014, Appli 17, Appli 1, Appli 17, Appli

NFORMATION FOR SEQ ID NO:

TELEPHONE: 202-508-9100

REGISTRATION NUMBER:

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                                                        COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

COMPUTER: IBM PC compatible

OBERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/047,041A

FILING DATE: 22-MAR-1993

CLASSIFICATION: 435

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 07/928,661

FILING DATE: 17-ANG-1992

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 07/446,584

APPLICATION NUMBER: US 07/446,584
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Patent No. 552767
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                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
ADDRESSEE: Banner & Allegretti, Ltd.
STREET: 1001 G Street, N.W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
LENGTH: 102 base pairs
TYPE: nucleic acid
            FILING DATE: 06-DEC-
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PUBLICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: Detec
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                                                                                                                                                                                                                                                                                                                                            CITY: Washington
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Fearon, Eric R.
Nigro, Janice M.
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                                               06-DEC-1989
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Pred. No. 3.8;
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Patent No.
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INFORMATION FOR SEQ ID NO:
                                          ATTORNEY/AGENT INFORMATION:
NAME: Kagan, Serah A
REGISTRATION NUMBER: 32141
REFERENCE/DOCKET NUMBER: 0110
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-508-9100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                      APPLICATION NUMBER: US/0
FILING DATE: 05-FEB-1997
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                 SOFTWARE: FastSEQ for Windows Version CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: NUMBER OF SEQUENCES: 3:
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ORIGINAL SOURCE:
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STREET: 1001 G Street, NW
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                                                                                                                                                                                                                                                                 OPERATING SYSTEM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               IITLE: A variation in the structure of the
ITLE: protein-coding region of the human p53 gene
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Brachmann,
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                                                                                                                                                                                                                                                                                  IBM Compatible
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              MUTATIONS WHICH SUPPRESS P53 CANCER MUTA- TIONS
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NUCLEIC ACIDS ENCODING P53
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22:
                                                                               01107.03170
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Pred. No.
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RESULT 5
US-08-472-239-1/c
; Sequence 1, Application US/08472239
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US-09-184-073-22/c
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                                                                                                                                                Query Match
Best Local Similarity
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Best Local :
                                                                                                                                                                                                                                                                                                                   ATTORNEY AGENT INFORMATION:
NAME: Kagan, Sarah A
REGISTRATION NUMBER: 32141
REFERENCE/DOCKET NUMBER: 0110
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-508-9100
                                                                                                                                                                                                                                                                INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    tent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: FASTSEQ for Windows Version 2.0 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/09/184,073
                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: Banner & Witcoff
STREET: 1001 G Street, NW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STREET: 1001
                                                                                                                                                                                                                   LENGTH: 133 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM CON
OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TLE OF INVENTION: NUCLEIC AC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            QUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDIUM TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY:
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                                                                                                                                   100.0%; Score 18; DB 100.0%; Pred. No. 3.8; tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%; Score 18; DB 2; Length 133; 100.0%; Pred. No. 3.8;
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NS WHICH SUPPRESS P53 CANCER MUTA- TIONS
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                                                           US-09-414-436-4
Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                        Sequence 4, Application US/09414436
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: GEORGE, Jr., Albert L.
APPLICANT: BHATNAGAR, Satish K.
APPLICANT: NAZARENKO, Irena
TITLE OF INVENTION: METHOD FOR ANALYZING A NUCLEOTIDE
TITLE OF INVENTION: SEQUENCE AND KIT THEREFOR
                                                                                                                                                                                                                                                                                                                                                                                                                09-414-436-4/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: (703) 836-20 INFORMATION FOR SEQ ID NO:
                                                                                                                                SOFTWARE: PatentIn Ver. 2.0 EQ ID NO 4
                                                                                                                                                                   EARLIER APPLICATION NUMBER: 60/103,849
EARLIER FILING DATE: 1999-10-09
NUMBER OF SEQ ID NOS: 5
                                                                          TYPE: DNA
ORGANISM: Homo
                                                                                                                                                                                                                                                                                                                                       APPLICANT: Dell'Acqua, Giorgio
                                                                                                                                                                                                                        TLE REFERENCE: p53f
URRENT APPLICATION NUMBER: US/09/414,436
URRENT FILING DATE: 1999-10-07
                                                                                                                                                                                                                                                                                                PPLICANT: Mann, Michael d
                                                                                                                                                                                                                                                                                                                                                                            tent No. 6294384
                                                                                                                ENGTH: 1070
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/472,239
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                              TLE OF INVENTION: Compositions and Methods Based Upon an Isoform of p53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
NAME: Mukai, Robert G.
REGISTRATION NUMBER: 28,
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES: 2
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TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         108
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P.O. Box 1404
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
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   100.0%;
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   Score 18;
Pred. No.
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   , DB 3;
. 4.3;
                   Length 1070;
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US-08-047-041A-13
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PSTEEMIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
CURRENT APPLICATION NUMBER: US/08/047,041A
                                                                                                                                                                                                                                                                                                      TELEFAX: 202-508-9299 (NFORMATION FOR SEQ ID NO: 13:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATA:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/928,661
APPLICATION TATE: 17-AUG-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE: 06-DEC-
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                       PUBLICATION INFORMATION:
AUTHORS: Harris, N.
JOURNAL: Mol. Cell. Biol
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE: 17-AUG-1992
PRIOR APPLICATION NUMBER: US 07/446,584
                                                                                                                                                HYPOTHETICAL: NO
ANTI-SENSE: NO
ORIGINAL SOURCE:
ORGANISM: Homo sapiens
                                                                                                                     POSITION IN GENOME:
                                                                                                                                                                                                                                                                                                                                           TELECOMMUNICATION INFORMATION: TELEPHONE: 202-508-9100
                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS LENGTH: 1303 base pair
                                                                                                                                                                                                                                                                                                                                                                                                                       TTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                TOPOLOGI.

JLECULE TYPE: CUN
                                                                                                                                                                                                                                                                                                                                                                         NAME: Kagan, Sarah A.
REGISTRATION NUMBER: 32,141
REFERENCE/DOCKET NUMBER: 01107.42917
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DDRESSEE:
                                                                                                                                                                                                                                              TRANDEDNESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NT: Fearon, Eric R.
NT: Nigro, Janice M.
F INVENTION: Detectic
F INVENTION: Gene
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                             4650-4656
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1001 G Street, N.W.
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Best Local S
Matches 18
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                   MOLECULE TYPE:
HYPOTHETICAL: N
ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/047,041A
FILING DATE: 22-MAR-1993
        MAP POSITION: 17p13.1
PUBLICATION INFORMATION:
AUTHORS: Harris, N.
TITLE: Molecular basis for heterogeneity of the
TITLE: human p53 protein
JOURNAL: Mol. Cell. Biol.
                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: 06-DEC-
PRIOR APPLICATION DATA:
APPLICATION NUMBER: U
FILING DATE: 29-MAR-1
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APPLICATION NUMBER:
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APPLICATION NUMBER: (
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                                                                                                      POSITION IN GENOME:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                               STRANDEDNESS:
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o. 5527676
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TER READABLE FORM:
TUM TYPE: Floppy disk
PUTER: IBM PC compatible
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                                                                                                                                                                                                                               nucleic acid
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1001 G Street, N.W.
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Widro, Janice M.
                                                                                                                                                                                                 linear
                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                           29-MAR-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   17-AUG-1992
                                                                                                                                                                                    CDNA
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                                                                                                                                                                                                                  double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Detection of Loss of the Wild-Type p53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US 07/446,584
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RESULT 10
US-08-347-792-1/c
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US-07-912-011-1/c
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US-08-047-041A-12
             Sequence 1, Application US/08347792
Patent No. 5573925
GENERAL INFORMATION: Thanos I
APPLICANT: Halazonetis, Thanos I
TITLE OF INVENTION: p53 Protein
TITLE OF INVENTION: Tetramerizat
                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                         Query Match
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Patent No. 5382510
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                   TELEPAX: 212-645-2054
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US/07/912
FILING DATE: 10-JUL-1992
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Feit, Irving N.
REGISTRATION NUMBER: 28,601
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Flopy disk
COMPUTER: ISM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                tent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: LE
TELECOMMUNICATION INFORMATION:
TELEPHONE: 212-645-1405
                                                                                                                                                                                                                                                                                                                      MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
ADDRESSEE: Imclone Systems Incorporated
STREET: 180 Varick Street
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PAGES: 401
                                                                                                                                                                                                                                                                                                                                     TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                        Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                     ENGTH:
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                                                                                                                                                                                                                                                      Similarity
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INVENTION: Probes for Detecting Mutant p53
                                                                                                                                                                        CCCGGAAGGCAGTCTGGC 110
                                                                                                                                                                                                      CCCGGAAGGCAGTCTGGC 18
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                                                                                                                                                                                                                                   100.0%; Score 18; 100.0%; Pred. No. tive 0; Mismatch
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          tis, Thanos D.
p53 Proteins With Altered
Tetramerization Domains
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US-08-431-357-1/c
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Patent No.
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Matches
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INFORMATION FOR SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM FO COMPATIBLE
COPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                             CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/347,792
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION: 530
ATTORNEY/AGENT INFORMATION:
NAME: Bak, Mary E.
REGISTRATION NUMBER: 31,2
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Halazonetis, Thanos D.
TITLE OF INVENTION: p53 Proteins With Altered
TITLE OF INVENTION: Tetramerization Domains
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: linear MOLECULE TYPE: DNA (genomic)
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 3
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ADDRESSEE: Howson an
TTORNEY/AGENT INFORMATION
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LOCATION:
                                                                                                                                                                                                                                                                         STREET: Spring House Corporate Cntr., PO Box 457 CITY: Spring House STATE: Pennsylvania
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                                                                                         FILING DATE:
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                                                                                                            APPLICATION NUMBER:
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5. 5721340
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Spring House
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                 28-NOV-1
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US-08-697-221-1
                                                                                                              TELECUMAN. 215-40-5818
TELEPAX: 215-40-5818
TELEFAX: 215-40-5818
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
SEQUENT: 1317 base pairs
TONGTH: 1317 base pairs
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                                                                                                                                                                                                        APPLICATION NUMBER: US/08/697,221
FILING DATE:
CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/004,802
FILING DATE: 22-SEP-1995
ATTORNEY/AGENT INFORMATION:
NAME: Kodroff, Cathy A.
REFERENCE/DOCKET NUMBER: 3980
REFERENCE/DOCKET NUMBER: WST64AUSA
TELECOMMUNICATION INFORMATION:
TELEPHONE: 215-540-9206
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
APPLICANT: Halazonetis, Thanc
TITLE OF INVENTION: Modified
TITLE OF INVENTION: Therefor
NUMBER OF SEQUENCES: 33
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INFORMATION FOR SEQ ID NO: 1:
                                                   MOLECULE TYPE:
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
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REFERENCE/DOCKET NUMBER: WST
TELECOMMUNICATION INFORMATION:
TELEPHONE: 215-540-9206
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STREET: Spring House Corporate Cntr., PO Box 457
CITY: Spring House
STATE: Pennsylvania
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local
                                                                                    TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY:
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STRANDEDNESS: double
                                  NAME/KEY:
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                                                                  DNA (genomic)
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                                                                                                                               RESULT 14
US-08-894-327-1/c
                                                                                                                                                                                                                                                                                                                                                  ; MOLECULE TYPE: cDNA
; ORIGINAL SOURCE:
; ORGANISM: Homo sapiens
US-08-392-542-1
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US-08-392-542-1/c
GENERAL INFORMATION:
APPLICANT: Halazonetis, Thanos
APPLICANT: Hartwig, Wolfgang
TITLE OF INVENTION: Peptides and peptidomimetics with
TITLE OF INVENTION: structural similarity to human p53 that activate
                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
Matches 18; Conserv
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Patent No. 6169073
                                                                                          Sequence 1, Application US/08894327 Patent No. 6245886
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Best Local Similarity
Matches 18; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: 202 508-9299
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION:
NAME: POSOTSKE, LAUTENCE H.
REGISTRATION NUMBER: 34,698
REFERENCE/DOCKET NUMBER: 0486
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202 508-9100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
CURRENT APPLICATION DATA:
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APPLICANT: Hartwig, Wolfgang
TITLE OF INVENTION: Peptides nad Peptidomimetics with
TITLE OF INVENTION: Structural Similarity to Human p53 That
TITLE OF INVENTION: Function
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/08/392,542
FILING DATE:
CLASSIFICATION: 514
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                                                                                                                                                                                                      127 CCCGGAAGGCAGTCTGGC 110
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   nucleic acid

DEDNESS: double
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District of Columbia
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1001 G Street, N.W.
                                                                                                                                                                                                                                                                         100.0%; Score 18; DB ilarity 100.0%; Pred. No. 4.3; Conservative 0; Mismatches
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Pred. No. 4.3;
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                                                                                                                                                                                                                                                                                                              DB 3;
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TITLE OF INVENTION: function

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; ORGANISM: Homo sapiens
US-08-894-327-1
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EARLIER FILING DATE: 1996-02-16
EARLIER APPLICATION NUMBER: 08/392,542
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URRENT APPLICATION NUMBER: US/08/894,327
URRENT FILING DATE: 1997-12-04
              NAME: Maebius, Stephen B.
REGISTRATION NUMBER: 35,264
REFERENCE/DOCKET NUMBER: 84
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 672-5300
                                                                                                                                                                                                                                                    FILING DATE: 18-MAY-1999
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: WO PCT/JP97/03019
FILING DATE: 29-AUG-1997
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/147,75
FILING DATE: 18-MAY-1999
                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: OBATA, Kazuaki
TITLE OF INVENTION: METHODS FOR TARGETING, ENRICHING
TITLE OF INVENTION: DETECTING AND/OR ISOLATING TARGET
TITLE OF INVENTION: USING RECA-LIKE RECOMBINASE
                                                                                                               FILING DATE: 29-AUG-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                       PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
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Local Similarity 100.0%;
                                                                                                                                                                                              APPLICATION NUMBER: JP 8 FILING DATE: 26-DEC-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE:
                                                                                                                                                       APPLICATION NUMBER: JP 8/229061
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       KIGAWA, Koji
YAMANAKA, Mikayo
KUSUMI, Kayo
MUXAI, Eli
OBATA, Kazuaki
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672-5399
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Pred. No.
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US-09-685-027-1/c
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                                                                                                                                                                                                                                                                                      Patent No. 64zuxxc
                                                                                                                                                                                                                                                                                                           Sequence 1, Approximately No. 6420118
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CURRENT FILING DATE: 1999-05-05
EARLIER APPLICATION NUMBER: 60/084,839
EARLIER FILING DATE: 1998-05-08
NUMBER OF SEQ ID NOS: 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Halazonetis, Thanos D.
APPLICANT: Stavridi, Elena S.
FITLE OF INVENTION: Modified p53 Tetramerization Domains Having
FITLE OF INVENTION: Amino Acid Substitutions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: human FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME/KEY: CDS
LOCATION: (136)..(1314)
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                                                                                                                    NUMBER OF SEQUENCE::

CORRESPONDENCE ADDRESS:

ADDRESSEE: Banner, Birch, McK
ATTRET: 1001 G Street, N.W.

Therefore D.C.

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               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                 Hartwig, TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                    APPLICANT: Halazonetis,
                                                                                                                                                                                                                                                                                                                                                                                                                       127
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5. 6388062
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                                                                                                                CITY: Washington, D.C.
STATE: District of Columbia
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PatentIn Release #1.0,
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                                                                                                                                                                                                            Wolfgang
Peptides nad Peptidomimetics with
Structural Similarity to Human p53 That Activate p53
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Pred. No. 4.3;
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Pred. No. 4
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                                                                                                                                                                  McKie & Beckett
   Version #1.25
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PCT-US95-15353-1/c
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                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US95/15353
                                       PRIOR APPLICATION NUMBER: US 08/431,357
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 00///
FILING DATE: 07 TYPORNEW //
                                                                                                               FILING DATE: 28-NOV-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 01
FILING DATE: 28-APR-1995
                                                                                                                                                                                          CLASSIFICATION:
PRIOR APPLICATION NUMBER:
                                     ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PLICANT: Halazonetis, Thanos D.
TLE OF INVENTION: p53 Proteins With Altered
TLE OF INVENTION: Tetramerization Domains
                                                                                                                                                                                                                                                      FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: 08/392,542
FILING DATE: <Unknown>
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: Homo sapiens SEQUENCE DESCRIPTION: SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: linear MOLECULE TYPE: cDNA
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TELEPHONE: 202 508-9100
TELEFAX: 202 508-929
WATION FOR SEQ ID NO: 1:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORIGINAL SOURCE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   18;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                            I: Spring House Corporate Cntr., PO Box 457
Spring House
: Pennsylvania
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REGISTRATION NUMBER: 34,698
REFERENCE/DOCKET NUMBER: 0486.48439
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/09/685,027 FILING DATE: 10-Oct-2000 CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: nucleic acid
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ilarity 100.0%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                  USA
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31,215
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-404-879A-332
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US-09-404-879A-332
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US-09-702-705-1634/c
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Best Local S
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GENERAL INFORMATIC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT FILING DATE: 1999-09-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT FILING DATE: 1999-09-24
NUMBER OF SEQ ID NOS: 393
                                                                                                                                                                                   quence 1634, Application US/09702705
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TILE OF INVENTION: COMPOS
TILE OF INVENTION: DIAGNO
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LOCATION:
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                                                                                                                                                                                                                                                                                                                                 Local Similarity es 16; Conserv
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                                                                                                                                                                                                                                                                           112 CCAGAAGGCAGTCTGGC 128
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           127 CCCGGAAGGCAGTCTGGC 110
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INVENTION: COMPOSITIONS AND METHODS INVENTION: DIAGNOSIS OF LUNG CANCER RENCE: 210121.478C14
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                                                   Fanger, Gary
Vedvick, Tom
Carter, Darrick
Retter, Marc
Mannion, Jane
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                                                                                                                            Wang, Tongtong
Bangur, Chaitanya S.
Lodes, Michael A.
                                            Fan, Liqun
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                                                                                                                                                                                                                                                                                                                                              94.1%;
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100.0%; Pred. No.
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Pred. No. 7
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                              FOR THE THERAPY AND
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RESULT 22
US-09-614-124B-1634/c
; Sequence 1634, Application US/09614124B
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-736-457-1634
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; ORGANISM: Homo sapiens
US-09-702-705-1634
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SEQ ID NO 1634
LENGTH: 447
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APPLICANT: Wang, Aijun
APPLICANT: Wang, Aijun
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF LUNG CANCER
FILE REFERENCE: 210121.478C15
CURRENT APPLICATION NUMBER: US/09/736,457
CURRENT FILING DATE: 2000-12-13
NUMBER OF SEQ ID NOS: 1864
SOFTMARE: FastSEQ for Windows Version 3.0
SEQ ID NO 1634
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atent No. 6509448
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                                                                                                                               APPLICANT: Wang, Tongtong
APPLICANT: Bangur, Chaite
APPLICANT: Lodes, Michael
APPLICANT: Fanger, Gary
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                                                                                                   APPLICANT:
                                                                       APPLICANT:
CURRENT FILING DATE:
                                      APPLICANT: Mannion, Jane
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR
TITLE OF INVENTION: DIAGNOSIS OF LUNG CANCER
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les 16; Conserv
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          APPLICATION NUMBER: US/09/614,124B
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                                                                             Bangur, Chaitanya S.
Lodes, Michael A.
Fanger, Gary
Vedvick, Tom
Carter, Darrick
Retter, Marc
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Retter, Marc
Mannion, Jane
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Lodes, Michael A.
Fanger, Gary
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r, Chaitanya
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94.1%;
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2001-07-11
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94.1%;
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Pred. No. 78;
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Pred. No. 78;
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                                                       THERAPY AND
                                                                                                                                                                                                                                                                                                                                                                            Length 447;
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GENERAL INFORMATION:
APPLICANT: ENDEGE, WILSON O., ET AL.
APPLICANT: ENDEGE, WILSON O., ET AL.
TITLE OF INVENTION: NOVEL HUMAN GENES AND GENE EXPRESSION
TITLE OF INVENTION: PRODUCTS: II
FILE REFERENCE: CCDNA-260XX
CURRENT APPLICATION NUMBER: U9/09/385,982
CURRENT FILING DATE: 1999-08-30
EARLIER APPLICATION NUMBER: 09/328,111
EARLIER FILING DATE: 1999-06-08
EARLIER FILING DATE: 1999-01-27
EARLIER APPLICATION NUMBER: 60/117,393
EARLIER APPLICATION NUMBER: 60/098,639
EARLIER APPLICATION NUMBER: 60/098,639
EARLIER FILING DATE: 1998-08-31
NUMBER OF SEQ ID NOS: 544
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SEQ ID NO 1634
LENGTH: 447
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SOFTWARE: FastSEQ for Windo
SEQ ID NO 1634
LENGTH: 447
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Patent No. 6667154
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Best Local S
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Best Local Similarity
SEQ ID NO 206
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TITLE OF INVENTION: COMPOSITIONS AND METI
TITLE OF INVENTION: DIAGNOSIS OF LUNG CAI
FILE REFERENCE: 210121.478C12
CURRENT APPLICATION NUMBER: US/09/671,325
CURRENT FILING DATE: 2000-09-26
NUMBER OF SEQ ID NOS: 1825
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16; Conserv
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                FastSEQ for Windows Version 3.0
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Vedvick, Tom
Carter, Darrick
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Mannion, Jane
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Bangur, Chaitanya
Lodes, Michael A.
                                                                                                                                                                                                                                                                                              Application US/09385982
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94.1%;
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94.1%;
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Pred. No. 78;
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Pred. No. 78;
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; LENGTH: 647
; TYPE: DNA
; ORGANISM: Human
US-09-222-575-81
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                                                                                                                                                                                                               GENERAL INFORMATION:
APPLICANT: Yuqui, Jiang
APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: XI, Jiangchun
TITLE OF INVENTION: COMPOSITIONS FOR THE TREATMENT
TITLE OF INVENTION: DIAGNOSIS OF BREAST CANCER AND
FILE REFERENCE: 210121.470C3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Yuqiu, Jiang
APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Xu, Jiangchun
TITLE OF INVENTION: Compositions for the Treatment and Diagnosis of Breast
TITLE OF INVENTION: and Methods for Their Use
FILE REFERENCE: 210121,470
CURRENT APPLICATION NUMBER: US/09/222,575
CURRENT FILING DATE: 1998-12-28
NUMBER OF SEQ ID NOS: 174
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 81
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 81, Application US/09222575 Patent No. 6387697
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Query Match
Best Local Similarity
                                                       TYPE: DNA
ORGANISM: Homo sapien
-09-389-681-81
                                                                                                                                                                                                                                                                                                                                                               Sequence 81, App
Patent No. 65182
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ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: misc feature
LOCATION: (1)...(633)
OTHER INFORMATION: n = A,T,C o:
                                                                                                                                                               CURRENT APPLICATION NUMBER: US/09/389,681A
CURRENT FILING DATE: 1999-09-02
NUMBER OF SEQ ID NOS: 463
                                                                                                          NUMBER OF SEQ ID NOS: 463
SOFTWARE: FASTSEQ for Windows Version 3.0
SEQ ID NO 81
LENGTH: 647
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94.1%;
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94.1%; Pred. No. 7
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Pred. No. 79;
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     Score 15.4; DB 4; Pred. No. 79;
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                    Length 647;
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APPLICANT: MUQIN, Jiang
APPLICANT: MUQIN, Jiang
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: MITCHAM, Jennifer L.
APPLICANT: MITCHAM, JENNIFER FOR THE TREATMENT AND
TITLE OF INVENTION: COMPOSITIONS FOR THE TREATMENT AND
TITLE OF INVENTION: DIAGNOSIS OF BREAST CANCER AND MET
FILE REFERENCE: 210121.470C2
FURLENT APPLICATION NUMBER: US/09/339,338A
CURRENT APPLICATION NUMBER: US/09/339,338A
CURRENT FILING DATE: 1999-06-23
NUMBER OF SEQ ID NOS: 315
SOFTMARE: FASTSEQ for Windows Version 3.0
SEQ ID NO 81
LENGTH: 647
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US-09-339-338-81
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APPLICANT: Jiang, Yuqiu
APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Xu, Jiangchun
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Best Local Similarity
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CURRENT APPLICATION NUMBER: US/09/620,405B
CURRENT FILING DATE: 2000-07-20
NUMBER OF SEQ ID NOS: 495
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Harlocker Susan L.
APPLICANT: Hepler, William T.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR
TITLE OF INVENTION: DIAGNOSIS OF BREAST CANCER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: DNA ORGANISM: Homo sapien
                                                                                                                                                                    TYPE: DNA
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Local Similarity 94.1%;
les 16; Conservative
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575
                                                                 16;
                              2 CCGGAAGGCAGTCTGGC 18
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CCAGAAGGCAGTCTGGC
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                                                                  Conservative
                                                                                 85.6%;
94.1%;
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METHODS FOR THEIR USE
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RESULT 29 US-09-433-826B-81

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                                                                                        Sequence 81, Application US/09285480 Patent No. 6590076
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                                                                         GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                 SEQ ID NO 81
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APPLICANT: Xu, Jiangchun
TITLE OF INVENTION: COMPOSITIONS FOR THE TREATMENT AND
                        APPLICANT: Yuqui, Jiang
APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jennifer L.
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                                                                                                                                                                                                                                                                                                                 LENGTH: 647
TYPE: DNA
ORGANISM: Homo sapien
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TYPE: DNA
ORGANISM: Homo sapien
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ITLE OF INVENTION: COMPOSITIONS FOR THE TREATMENT AND ITLE OF INVENTION: DIAGNOSIS OF BREAST CANCER AND METHODS FOR THEIR USE ILB REFERENCE: 210121.470C4
                                                                                                                                                                                                                                                                                                                                                                                                                                                   LICANT: Harlocker, Susan L.
LICANT: Hepler, William T.
LE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
LE OF INVENTION: DIAGNOSIS OF BREAST CANCER
                                                                                                                                                                                                                                                       Local
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Mitcham, Jennifer L.
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94.1%;
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Pred. No. 79;
0; Mismatches
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0; Mismatches
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Pred. No. 79
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US-08-447-179-7/c
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SEQ ID NO 81
LENGTH: 647
TYPE: DNA
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Best Local Similarity
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CURRENT FILING DATE: 2001-C
NUMBER OF SEQ ID NOS: 547
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/09/285,480 CURRENT FILING DATE: 1999-04-02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: DIAGNOSIS OF BREAST CANCER AND METHODS FOR THEIR USE FILE REFERENCE: 210121.470C1
                                                                                                                                                                                equence 7, Application US/08447179 atent No. 5744303
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Homo sapien
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PPLICANT: Henderson, Robert A.
ITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
ITLE OF INVENTION: DIAGNOSIS OF BREAST CANCER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 647
                                                                                                                                      APPLICANT:
                                                       CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity
                             STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REFERENCE: 210121.470C9
                                         DDRESSEE:
                                                                                  E OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     575
                                                                                                                                                                                                                                                                   575 CCAGAAGGCAGTCTGGC 591
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                                                                                                                                                                    INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                          647
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     FastSEQ for Windows Version 3.0
Massachusetts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Harlocker, Susan L.
Hepler, William T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mitcham, Jennifer L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Application US/09834759
                             225 Franklin Street
                                                                                                                                      Friend, Stephen
                                                                                                                                                                                                                                                                                                                           Conservative
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                                                                                                          Frebourg, Thierry
Ishioka, Chikashi
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                                                                                                                                                        Richard
                                                                                   FUNCTIONAL
SUPPRESSOR
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94.1%;
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                                        Richardson
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                                                                                                                                                                                                                                                                                                                                       Score 15.4;
Pred. No. 79
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Pred. No. 7
                                                                                   ASSAY
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                                                                                               FOR TUMOR
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                                                                                                                                                                                                                                                                                                                                                    Length 647;
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US-09-445-247-21/c
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FILING DATE:
FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/046,033
FILING DATE: 12 APRIL 1993
APPLICATION NUMBER: 07/956,696
FILING DATE: 10 OCTOBER 1992
ATTORNEY/AGENT INFORMATION:
NAME: Clark, Paul T.
NAME: Clark, Paul T.
00786/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 21, Application Patent No. 6410238
GENERAL INFORMATION:
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TELEX: 200154
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 542-5070
TELEPAX: (617) 542-8906
TELEPAX: (617) 542-8906
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PS/2 Model 50Z or 55SX OPERATING SYSTEM: MS-DOS (Version 5. SOFTWARE: Wordperfect (Version 5.1) CURRENT APPLICATION DATA:
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                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION UNMERS: US/09/445,247
APPLICATION UNMERS: US/09/445,247
FILING DATE: 03-Dec-1999
CLASSIFICATION: <Unknown>
PRIOR APPLICATION OMMBER: US/08/870,126
APPLICATION NUMBER: US/08/870,126
FILING DATE: 06-UN-1997
FILING DATE: 06-UN-1997
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STRANDEDNESS: singl
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                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:

ADDRESSEE: Howson and Howson
STREET: Spring House Corporate Cntr, P O Box 457
CITY: Spring House
STATE: Pennsylvania
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Wistar Institute of Anatomy & Biology Prendergast, George C.
                                       ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: Box-Dependent MYC-Interacting Protein
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NAME: Bak, Mary E.
REGISTRATION NUMBER: 31,215
                                                                                                                                                                                                                                                                                                                 COUNTRY: USA
ZIP: 19477
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Application US/09445247
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RESULT 35
US-08-094-071-1/c
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARS: PATENTION DATA:
APPLICATION DATA:
APPLICATION UMBER: US/08/094,071
FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 719,172
FILING DATE: 21-UWN-1991
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
APPLICANT: Reynolds, Frederick H.
APPLICANT: Sorvillo, John M.
APPLICANT: Zeheb, Ron J.
APPLICANT: Stephenson, John R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEPHONE: 215-540-9200
TELEPAX: 215-540-5818
INFORMATION FOR SEQ ID NO: 21:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                        APPLICATION NUMBER: 1
FILING DATE: 18-JAN-
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US 2: FILING DATE: 17-JAN-1989 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US 6: FILING DATE: 01-FEB-1991 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Stephenson, TITLE OF INVENTION: Imm TITLE OF INVENTION: mut
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                                                                                                             APPLICATION NUMBER: FILING DATE: 23-JUL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: WST60DPCT TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FEATURE:
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STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FOPOLOGY: unknown
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                                                                                                                                                                                                                              TUMBER: US 298,837
18-JAN-1989
MBER: US 767,862
21-AUG-1985
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biological fluids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US 649,566
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                                                                                                                                                  US 885,627
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Pred. No. 1
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Matches
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Best Local
                                                                                                                                                    GENERAL INFORMATION:
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                                                                                                                                                                                    -US92-00878-1/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PUBLICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NTI-SENSE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELECOMMUNICATION INFORMATION: TELEPHONE: (212) 977-9550 TELEFAX: (212) 664-0525
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
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PRIOR APPLICATION DATA:
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APPLICATION NUMBER:
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                                             TUMBER OF SEQUENCES:
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                             ORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: AU 6
FILING DATE: 18-AUG-1986
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: NZ 217,209
FILING DATE: 14-AUG-1986
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER:
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                DDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TRANDEDNESS: single
                                                     LE OF INVENTION: Immunoassay for Detection of mutant p53
LE OF INVENTION: polypeptide in biological fluids
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                                                                                                                                                                                                                                             206 CCGGAAAGGCAGTCTGGC 189
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                                                                                                                                                                                                                                                                                                                     Similarity
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Raiston, R.
Helfman, D. M.
Adams, T. E.
Adams, T. Cloning and In-Vitro
Molecular Cloning and Councillation for Human
Transion of a C-DNA Clone for Human
Transion p53
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DATE: 20-AUG-1986
                                                                                                                                                               Application PC/TUS9200878
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               activated p53 oncogene
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E: Cooper and Dunham
30 Rockefeller Plaza
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Williamson, N. M.
                                                                                                                                                                                                                                                                                                       Conservative
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                                                                                                                  Reynolds, Frederick H. Sorvillo, John M.
                                                                                     Stephenson, John R.
                                                                                                       Zeheb, Ron J
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Pred. No. 1.7e.
0; Mismatches
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STATE: New York
COUNTRY: USA
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TELEX: 422523 coop ui INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 1757 base pairs PRIOR APPLICATION UNMEER: US 885,627
APPLICATION NUMBER: US 885,627 PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 719,172
FILING DATE: 21-JUN-1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 649,566
APPLICATION NUMBER: US 649,566 COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.2
CURRENT APPLICATION DATA: PUBLICATION INFORMATION: PRIOR APPLICATION DATA: COMPUTER READABLE FORM: MEDIUM TYPE: Floppy TELECOMMUNICATION INFORMATION: ATTORNEY/AGENT INFORMATION: PRIOR APPLICATION DATA: PRIOR APPLICATION DATA:
APPLICATION NUMBER: MOLECULE TYPE: C MMEDIATE SOURCE: FILING DATE: 18-AIC REFERENCE/DOCKET NUMBER: 28 APPLICATION NUMBER: FILING DATE: 19920 STRANDEDNESS: APPLICATION NUMBER: APPLICATION NUMBER: APPLICATION NUMBER: CLASSIFICATION: NUCLEIC ACID Molecular Cloning and In-Vitro Expression of a C-DNA Clone for Human Cellular Tumor Antigen P53 Harlow, E.
Williamson, N. M.
Ralston, R.
Helfman, D. M. activated p53 oncogene CATION DATA: ION NUMBER: CA 516,260 ATE: 19-AUG-1986 CATION DATA: ON NUMBER: ON NUMBER CATION DATA: CATION DATA: CATION DATA: linear John P. 28,678 Floppy disk UMBER: AU 61540/86 18-AUG-1986 01-FEB-1991 14-AUG-1986 19920131 CDNA single MBER: JP 195121/86 20-AUG-1986 MBER: EP 86.111 527.7 20-AUG-1986 21-AUG-1985 18-JAN-1989 17-JAN-1989 664-0525 977-9550 US 298,776 NZ 217,209 US 767,862 Release #1.24 US 298,837 PCT/US92/00878 ۳. 23384-в

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US-08-870-126-12/c
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Patent No. 6048702
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Best Local Similarity
Matches 16; Conserv
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: FIRM PC-DOS/MS-DOS
COMPUTER: FORM: PC-DOS/MS-DOS
COMPUTER: FORM: PC-DOS/MS-DOS
COMPUTER: PC-DOS/MS-D
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NFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 8078 base pair
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 215-540-9200
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APPLICATION NUMBER: 1
FILING DATE: 05-MAY-
PRIOR APPLICATION DATA:
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SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US 08/652,972 FILING DATE: 24-MAY-1996 ATTORNEY/AGENT INFORMATION:
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                                                                                             NAME/KEY: exon
LOCATION: 2409..2432
OTHER INFORMATION: /:
                                                                                                                                                                                                                                     NAME/KEY: exon
LOCATION: 1841..1948
OTHER INFORMATION: //
                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: nucleic acid
STRANDEDNESS: double
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TLE OF INVENTION: MYC-Interacting Protein (Bin1) and Uses Therefor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STREET: Spring House Corporate Cntr, P O Box 457 CITY: Spring House
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exon
3308..3601
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Pred. No. 1
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US-09-445-247-12/c
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                                                                                                                                                                                              TELEFAX: 215-540-5818
INFORMATION FOR SEQ ID NO: 12:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:

NAME: Bak, Mary E.

REGISTRATION NUMBER: 31,215

REFERENCE/DOCKET NUMBER: WST60DPCT

TELEPHONE: 215-540-9200

TELEPHONE: 215-540-9200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION NUMBER: US/09/445,247

APPLICATION NUMBER: US/09/445,247
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CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
                            MOLECULE TYPE: DNA (genomic)
                                                                                             TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US 08/870,126 FILING DATE: 06-JUN-1997
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                                                                                                                                                              ENGTH: 8078 base pairs
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Sequence 12, Application Patent No. 6410238
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity 88.9%;
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LOCATION: 5334..5435
OTHER INFORMATION: //
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LOCATION: 4944..5054
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                                                                                    TITLE OF INVENTION: Box-Dependent MYC-Interacting Protein (Bin1) Compositions and Uses Therefor
                                                                                                                                         APPLICANT: Wistar Institute of Anatomy & Biology Prendergast, George C.
                                                                         NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                /KEY: exon
FION: 7223..7783
R INFORMATION: /n
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 STREET: Spring House
CITY: Spring House
NDONDENCE THOUSEN and Howson
Thouse Corporate Cntr, P O Box 457
                                                                                                                                                                                                                Application US/09445247
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4613..4702
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Pred. No. 1.8e
0; Mismatches
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US-08-652-972A-6/c
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; OTHER INFORMATION: /note= "exon 16"
; SEQUENCE DESCRIPTION: SEQ ID NO: 12:
US-09-445-247-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local :
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                                                        APPLICATION NUMBER: US/08/652,972A
PILING DATE: 24-MAY-1996
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/435,454
FILING DATE: 05-MAY-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     quence 6, Application US/08652972A
tent No. 5723581
                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Prendergast, George C.

APPLICANT: Sakamuro, Daitoku

TITLE OF INVENTION: Murine and Human Box-Dependent

TITLE OF INVENTION: MYC-Interacting Protein (BIN1) and Uses Therefor
                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IRM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                       SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
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FELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                   ADDRESSEE: Howson and Howson STREET: Spring House Corporate CITY: Spring House STATE: Pennsylvania
            NAME: Bak, Mary E.
REGISTRATION NUMBER: 31,215
REFERENCE/DOCKET NUMBER: WS
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Local Similarity 88.9%;
es 16; Conservativo
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LOCATION:
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LOCATION: 4944..5054
OTHER INFORMATION: /I
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LOCATION: 4613..4'
OTHER INFORMATION:
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LOCATION: 3494..3601
CTHER INFORMATION: /note=
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LOCATION:
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OTHER INFORMATION:
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LOCATION: 2409..2432
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Pred. No. 1.8e+02;
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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NAME/KEY: exon
11851..12240
118710N: /note=
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LOCATION: 14130...
OTHER INFORMATION:
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LOCATION: 11519..11850
OTHER INFORMATION: /no
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LOCATION: 4441..11518
OTHER INFORMATION; /note=
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NAME/KEY: exon
175..4365
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LOCATION: 325..1618
OTHER INFORMATION:
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LOCATION: 3225
OTHER INFORMATION:
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LOCATION: 1619..31
OTHER INFORMATION:
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LOCATION: 1..324
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OTHER INFORMATION: /note=
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10738 CCCGGAAGGCAGGCTGCC 10721
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                                            ch 82.2%; 1 Similarity 88.9%; 16; Conservative (
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DEDNESS: double
                     CCCGGAAGGCAGTCTGGC 18
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14130..14985
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                                                                  Length 14985;
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PCT-US96-06231A-6/c
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VEORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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APPLICATION NUMBER: US 0
FILING DATE: 05-MAY-1995
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TITLE OF INVENTION: Murine and Human Box-Dependent
TITLE OF INVENTION: Myc-Interacting Protein (BIN1)
                                                                                                                                                                       OTHER INFORMATION: FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: PatentIn Release #1.0, Version CURRENT APPLICATION DATA:
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RELECOMMUNICATION INFORMATION:
TELEPHONE: 215-540-9200
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                                                                                                                                                                                                                                                              NAME/KEY:
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LOCATION:
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LOCATION: 325..1618
OTHER INFORMATION:
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LOCATION: 7209
OTHER INFORMATION:
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LOCATION: 3175..43
OTHER INFORMATION:
                                 NAME/KEY:
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TORNEY/AGENT INFORMATION:
                                                                OTHER INFORMATION:
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Spring House Corporate Cntr, P O Box 457
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3175..4365
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US-09-103-840A-1/c
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                                                                                                                              Query Match
Best Local S
Matches 16
Sequence 1, Application US/09103840A Patent No. 6294328
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                     FILE REFERENCE: 24366-ZUUU/...V
CURRENT APPLICATION NUMBER: US/09/103,840A
                                                                                                                                                                                                                                                                                                                    CURRENT FILING DATE: 1: NUMBER OF SEQ ID NOS: 2
                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: DNA SEQUENCES FOR STRAIN ANALYSIS IN MYCOBACTERIUM TITLE OF INVENTION: TUBERCULOSIS
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                                                                                                                                                                                             OTHER INFORMATION: CDC 1551
OTHER INFORMATION: "n" bases at various positions throughout the sequence OTHER INFORMATION: represent a, t, c or g
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   tent No.
                                                                                                                                                                                                                                                                                                                                                                                                                             PPLICANT:
                                                                                                                                                                                                                                                    ORGANISM: Mycobacterium tuberculosis
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                                                                             1092940 CCCGGACGGCAGGCTGGC 1092923
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les 16; Conserv
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OTHER INFORMATION: /no
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TION: 11851..12240
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12241..14129
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Pred. No. 1.9e
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                                                                                                                                                       Length 4403765;
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GENERAL

FLEISCHMAN, Robert D. WHITE, Owen R.

FRASER, Claire M.

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US-08-256-077-3
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                                                                                                                                                                                                                                                                                                                                                                                                            ; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: 148
; OTHER INFORMATION: n=a, g, c
US-09-621-976-11048
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 43
US-09-621-976-11048/c
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                                                                                                                                                                                       Sequence 3, Applic Patent No. 5654188
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                                                                                                                                                                                                                                                                                                                                               Matches
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                               Query Match
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SOFTWARE: Patent.pm
EQ ID NO 11048
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/621,976
CURRENT FILING DATE: 2000-07-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Giordano, J.Y.
TITLE OF INVENTION: ESTs and Encoded Human Proteins
FILE REFERENCE: GENSET.054PR2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILE REFERENCE: 24366-20007.00
CURRENT APPLICATION NUMBER: US/09/103,840A
CURRENT FILING DATE: 1998-06-24
NUMBER OF SEQ ID NOS: 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Dumas Milne Edwards, J.B. APPLICANT: Jobert, S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: PatentIn Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Mycobacterium tuberculosis
OTHER INFORMATION: H37Rv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 4411529
TYPE: DNA
                                                                                                    APPLICANT: Ellmeier, Wili
APPLICANT: Weith, Andreas
TITLE OF INVENTION: Neurol
NUMBER OF SEQUENCES: 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ENGTH: 252
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PPLICANT: VENTER, John C.
ITLE OF INVENTION: DNA SEQUENCES FOR STRAIN ANALYSIS IN MYCOBACTERIUM
ITLE OF INVENTION: TUBERCULOSIS
                                                                                         CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1092941 CCCGGACGGCAGGCTGGC 1092924
                                                              STREET:
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                                                                           DDRESSEE:
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16; Conserv
                                                                                                                                                                       INFORMATION:
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15; Conserva
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                                           Washington
20005-3934
                              D.C.
                                                                                                                                                                                                       Application US/08256077
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                                                            1100 New York Avenue,
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                                                                                                                        Neuroblastoma-Associated Regulator Gene
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93.8%;
                                                            Kessler, Goldstein & ork Avenue, Suite 600
                                                                                                                                                        Wilfried
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Pred. No. 2.3e
0; Mismatches
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0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 360 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: (202) 371-2540
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
NAME: ASMOOD, Robert W.
REGISTRATION NUMBER: 32,893
REFERENCE/DOCKET NUMBER: 06:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                    COUNTRY: U.S.A.
ZIP: 2005-3934
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM FC compatible
COMPUTER: IBM FC COMPUTER:
COMPUTER: PG-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                SOFTWARE: PatentIn Rel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
ADDRESSEE: Sterne, Kessler, Goldstein & Fox P.L.L.C.
STREET: 1100 New York Avenue, Suite 600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELECOMMUNICATION INFORMATION: TELEPHONE: (202) 371-2600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
MOLECULE TYPE:
                                                                                                                                                                                                                                              APPLICATION NUMBER: FILING DATE: 06-JUN CLASSIFICATION: 43
                                                                                                                                                                                                                                                                                                                                                                                                                                               STATE: D.C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                              STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/08/256,077 FILING DATE: 23-JUNE-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local
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                                                                                                                                         ELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LE OF INVENTION: Neuroblastoma-Associated Regulator Gene
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               l Similarity
15; Conserv
                                               nucleic acid
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                                                                                                                        (202) 371-2540
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Weith, Andreas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ellmeier, Willfried
                                                                                                                                         (202)
                                                                                                                                                                                                                                                                 06-JUN-1995
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                                                                                                                                         371-2600
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                                                                                                                                                                                                                                                                                   US/08/466,127
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Pred. No. 2.4e+02;
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; LOCATION:
US-08-466-127-3
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US-08-256-077-1
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                                                                                                                                                                                                                                                                                                                                                                                                NAME: Esmond, Robert W.
REGISTRATION NUMBER: 32,893
REFERENCE/DOCKET NUMBER: 0652
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 371-2600
TELEPAX: (202) 371-2540
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
                                                                  Sequence 1, Applic Patent No. 5683878
                                                                                                                                                                                                                                  Query Match 80.4
Best Local Similarity 93.4
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 1, Applic
Patent No. 5654188
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY: U.S.A.

COUNTRY: U.S.A.

ZIP: 20005-3934

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

COMPATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Ve

CURRENT APPLICATION DATA:
                                                  GENERAL INFORMATION:
APPLICANT: Ellmeier, Willfried APPLICANT: Weith, Andreas TITLE OF INVENTION: Neuroblast
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STREET: LL. STREET: Washington CITY: Washington CTATE: D.C.
                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE: cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Ellmeier, will Applicant: Weith, Andreas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/08/256,077
FILING DATE: 23-UUNE-1994
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: Neuroblastoma-Associated Regulator Gene
NUMBER OF SEQUENCES: 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local
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                                                                                                                                                                   176 CGGAACGCAGTCTGGC 191
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Similarity 93.8%;
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                                                                               Application US/08466127
                                                                                                                                                                                                 CGGAAGGCAGTCTGGC 18
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1100 New York Avenue,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
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Neuroblastoma-Associated Regulator Gene
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93.8%; Pred. No. 2.50
cive 0; Mismatches
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ork Avenue, Suite 600
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Pred. No. 2
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SOFTWARE: PERL Program
SEQ ID NO 7
LENGTH: 8137
TYPE: DNA
ORGANISM: Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; NAME/KEY:
; LOCATION:
US-08-466-127-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-09-566-921-7/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 93.0
                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                               Patent No.
                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/566,921
CURRENT FILING DATE: 2000-05-05
                                                                                                                                                                                                                                                                                                                                     APPLICANT: Loring, Jeanne F.
APPLICANT: Tingley, Debora W.
APPLICANT: Edwards, Carla M.
APPLICANT: Edwards, GENES EXPRESSED IN ALZHEIMER'S DISEASE
TITLE OF INVENTION: GENES EXPRESSED IN ALZHEIMER'S DISEASE
                                                                                                                                                                                                                                                                                    NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: (202) 371-25
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN PC-BOS/MS-DOS
SOFTWARE: PATENTIN DATA:
APPLICATION NUMBER: US/08/466,127
FILING DATE: 06-JUN-1995
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
                                                                                    NAME/KEY: unsure
LOCATION: 3488-3788
OTHER INFORMATION: a,
                                                                                                                                FEATURE:
NAME/KEY: misc feature
OTHER INFORMATION: Incyte ID No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY: lir
MOLECULE TYPE:
FEATURE:
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REGISTRATION NUMBER: 32,893
REFERENCE/DOCKET NUMBER: 06:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 371_2600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ZIP: 20005-3934
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: U.S.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                              INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      176 CGGAACGCAGTCTGGC 191
                                                                                                                                                                                                                                                                                                                                                                                                                                                            7, Application US/09566921
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3 CGGAAGGCAGTCTGGC 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Washington
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37..481
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77.8%; Score 14; DB 4; L
100.0%; Pred. No. 4.4e+02;
tive 0; Mismatches 0;
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Pred. No. 2.5e+02;
                                                                                       or other
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                                 Length 8137;
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US-08-721-986-1/c
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                                                                                               Patent No.
                                                                                                                                                                                                                                                            Query Match
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-08-721-986-1
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                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: 908-594-4720
NFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC pcs/MS-Dos
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APPLICATION NUMBER: US/08
FILING DATE: 08-APR-1994
ATTORNEY/AGENT INFORMATION:
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SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/721,986
FILING DATE:
     FITLE OF INVENTION:
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ADDRESSEE: JOSEPH A. COPPOLA
STREET: P.O. BOX 2000; 126 E. LINCOLN AVENUE
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LECOMMUNICATION INFORMATION:
TELEPHONE: 908-594-6734
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                                                                                                                                                                                                                                                                                                                                                                                      TYPE: nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION:
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                                                                                                                                                                                           103 CCCGGAAGGCAATCCGG 87
                                                                                  INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                       157 base pairs
                                                                                                                                                                                                                                                       76.7%; Score 13.8; ilarity 88.2%; Pred. No. 4.5 Conservative 0; Mismatches
                                                               NICHOLSON, DONALD W.
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                                                                                                               ication US/08225487A
                                                                                                                                                                                                                                                                                                                                                          linear
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                                                 AMBERDEEN
                                                                                                                                                                                                                                                                                                                                                                      single
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DNA ENCODING PRECURSOR OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DNA ENCODING PRECURSOR OF INTERLEUKIN-1BETA CONVERTING ENZYME -RELATED CYSTEINE PROTEINASE II (ICE rel-II)
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                                JOHN P.
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Search completed: March Job time : 98 secs
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Best Local (
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                                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION:
NAME: COPPOLA, JOSEPH A
REGISTRATION NUMBER: 39,413
REFERENCE/DOCKET NUMBER: 19
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
LENGTH: 157 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: Patentin Release #1.0, Version #1.30 CURRINT APPLICATION DATA: APPLICATION NUMBER: US/08/225,487A FILING DATE: 08-APR-1994 CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                           MOLECULE TYPE:
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ZIP: 07065
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                                                          103 CCCGGAAGGCAATCCGG 87
                                                                                                                                 Similarity
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C: P.O. BOX 2000; 126 E. LINCOLN AVENUE
RAHWAY
                                                                                     CCCGGAAGGCAGTCTGG 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NEW JERSEY
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                                                                                                                   Conservative
                                                                                                                                                                                                         linear
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                                                                                                                                                                                         DNA (genomic)
                                                                                                                                                                                                                      single
            5, 2004, 01:18:20
                                                                                                                               76.7%;
88.2%;
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                                                                                                                               Score 13.8;
Pred. No. 4.
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                                                                                                                   Mismatches
                                                                                                                               DB 3;
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VERSION
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BQ224113/c
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  88
                                                             Contact: Robert Strausberg, Ph.D.

Email: Gappbs-remail.nih.gov

Tissue Procurement: ATCC

CDNA Library Preparation: Life Technologies, Inc.

CDNA Library Arrayed by: The I.M.A.G.E. Consortium

CDNA Library Arrayed by: Genourt Bioscience Corporation

DNA Sequencing by: Agenourt Bioscience Corporation

Clone distribution: MGC clone distribution information can
                                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bass 1 to 818)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
Unpublished (1999)
  found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov
                                                                                                                                                                                                                                                                                                                                                                                                                BQ224113
BQ224113.1 GI:20405513
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Plate: LLAM14126 row: c column: 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
Unpublished (1999)
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BU902747
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                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                         5', mRNA sequence.
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AGENCOURT 7558930 NIH MGC 92 Homo sapiens
5'. mRNA Remierra
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Mammalia; Eutheria;
1 (bases 1 to 817)
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                                                                                                                                                                                                                                                                                                                                                                       sapiens (human)
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Location/Qualifiers
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/lab host="DH10B (phage-resistant)"
/clone_lib="NIH_MGC_71"
/clone_tib="NIH_MGC_71"
/note="Organ: uterus; Vector: pCMV-SPORT6; Site_1: NotI;
Site_2: SalI; Cloned unidirectionally. Primer: Oligo dT
Average insert size 2.1 kb. "
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /db_xref="taxon:9606"
/clone="IMAGE:6528131"
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Pred. No.
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VERSION
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BQ228424/c
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  Query Match
Best Local S
Matches 18
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NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
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BQ228424.1 GI:20409824
EST.
                                                                                                                                                                                                                                                                                                                                                                 High quality sequence stop: 611.
                                                                                                                                                                                                                                                                                                                                                                                  http://image.llnl.gov
Plate: LLAM13326 row: c column:
                                                                                                                                                                                                                                                                                                                                                                                                                           found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Email: cgapbs-r@mail.nih.gov
Tissue Procurement: DCTD/DTP/Gazdar
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
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Mammalia; Eutheria; Primates;
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AGENCOURT 7657440 NIH MGC 68 Homo sapiens cDNA
5'. mRNA TEOTICOTO
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                     Similarity
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100.0%; Score 18; llarity 100.0%; Pred. No. Conservative 0; Mismatch
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                                                                                                          /clone_lib="NIH_MGC 68"
/note="Organ: lung; Vector: pCMV-SPORT6; Site_1: l
/note="Organ: lung; Vector: pCMV-SPORT6; Site_1: l
Site_2: Sall; Cloned unidirectionally. Primer: Ol
Average insert size 1.8 kb. Library constructed
Technologies."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /tissue_type="embryonal carcinoma, cell line"
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                                                                                                                                                                                                                                            clone="IMAGE: 6059449"
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Pred. No. 6.1e+02;
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                     DB 13;
6.1e+02;
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AUTHORS
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VERSION
KEYWORDS
SOURCE
ORGANISM
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AU139600/c
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Search completed: March 5, 2004, 01:16:44
Job time : 2545 secs
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                                                                                                                                                        Query Match 100.0%; Score 18; DB 9; Length 847; Best Local Similarity 100.0%; Pred. No. 6.1e+02; Matches 18; Conservative 0; Mismatches 0; Indels
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HRI human cDNA project
Unpublished (2000)
Contact: Takao Isogai
Genomics Laboratory
Helix Research Institute
1537-3 Yana, Kisarazu, Chiba 292-0812, Japan
Tel: 81-438-52-3975
Fax: 81-438-52-3986
                                                                               669 CCCGGAAGGCAGTCTGGC 652
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                                                                                                                                                                                                                                                                                                                                                                                                     Email: genomics@hri.co.jp
HRI human cDNA project; 5'- & 3'-end one pass sequencing: Helix
Research Institute; cDNA library construction: Department of
Virology, Institute of Medical Science, University of Tokyo, and
Helix Research Institute.
Location/Qualifiers
1. .847
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Homo sapiens (buman)
Homo sapiens
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.

1 (bases 1 to 847)
Ota, T., Nishikawa, T., Suzuki, Y., Ishii, S., Saito, K., Kawai, Y.,
Yamamoto, J., Wakamatsu, A., Nakamura, Y., Nagai, T., Sugano, S. and
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AU139600
AU139600.1 GI:11001121
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